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COX-2 and PGE2-dependent immunomodulation in breast cancer

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Abstract

COX-derived prostanoids play multiple roles in inflammation and cancer. This review highlights research examining COX-2 and PGE₂-dependent regulation of immune cell polarization and function within the tumor microenvironment, particularly as it pertains to breast cancer. Appreciating PGE₂-mediated immunomodulation is important in understanding how tumors evade immune surveillance by reeducating infiltrating inflammatory and immune cells to support tumorigenesis. Elucidation of the multiple and complex influences exerted by tumor stromal components may lead to targeted therapies in breast and other cancers that restrain microenvironmental permissiveness and maintain natural defenses against malignancies.

Keywords

Breast cancer; COX-2; Macrophage polarization; Microenvironment; PGE2; Regulatory T lymphocytes; Myeloid-derived suppressor cells; Immune suppression

1. Introduction

Cyclooxygenase (COX) is the enzyme responsible for the conversion of arachidonic acid into the various prostanoids, a family of lipid mediators that have widespread and diverse biological function [1]. COX exists in two main isoforms, COX-1, which is predominantly constitutive and responsible for generation of prostanoids for "housekeeping functions", and COX-2, the inducible isoform, which contributes prostanoids involved in a variety of growth and inflammatory events [1,2]. Synthesis of eicosanoids begins after the release of arachidonic acid (AA) from membrane phospholipids through the action of cytosolic phospholipase A_2 . COX-1/COX-2, also known as prostaglandin G/H synthase 1/2, converts AA into prostaglandin (PG) G_2 and then reduces PG G_2 to PGH $_2$. PGH $_2$ can be metabolized by the various PG synthases into PGD $_2$, PGE $_2$, PGF $_2$ a, PGI $_2$, and thromboxane (TX) A_2 , which then act via distinct downstream G protein-coupled receptors.

A large body of work describing a link between inflammation and cancer [3] has generated intense interest in targeting COX enzymes, COX-2 in particular, for cancer therapy or chemoprevention. COX-2 is upregulated in 40% of breast cancers, with up to 84% increases in some studies [4]. Clinical studies have noted a reduced risk for breast, lung, prostate, and

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colon cancers after treatment with non-steroidal anti-inflammatory drugs (NSAIDs), which non-selectively inhibit COX-1 and COX-2, or with selective inhibition of COX-2 [5]. The beneficial effects of aspirin are less clear in part because many studies do not distinguish between consumption of low dose aspirin, whose effect is limited to inhibition of platelet COX-1 function, and higher doses that inhibit systemic function of both isozymes. In the Women's Health Initiative observational study, chronic regular use of NSAIDs was associated with reduced risk of breast cancer but subgroup analysis revealed no effect of low dose aspirin (<100 mg) [6]. Similarly, the Women's Health Study, a long term randomized trial, showed no effect of low dose aspirin every other day on breast cancer incidence [7]. Reduced risk of breast cancer death and distant recurrence, but not incidence of primary disease, was associated with regular aspirin use in the prospective observational Nurses' Health Study but dose was not reported [8]. In contrast, in another recent study, lifetime aspirin use was associated with a 32% decreased risk of breast cancer, though, again, no information on dosage was collected [9]. Analysis of eight aspirin trials revealed reduced cancer death that was independent of dose across several common cancers although scant information was available in breast cancer [10].

Certain COX-2-derived products, particularly PGE₂, are known to act via classical cancer signaling pathways in primary tumor cells to promote tumorigenesis. Recent evidence has shined a spotlight not only on the tumor cell itself, but the tumor microenvironment, or stroma, which surrounds the tumor. This is evidenced by Hanahan and Weinberg recently updating their landmark review of the hallmarks of cancer to include microenvironment specific components [11]. The microenvironment contains multiple resident and infiltrating cells, including immune cells, along with the growth factors and cytokines that they release. A supportive tumor microenvironment appears crucial for the development of a tumor as well as its transition to malignancy, and the characteristics of a pro-tumorigenic microenvironment has been well reviewed [12]. This review will focus on tumor evasion of immune surveillance, and how COX-2-derived PGE₂ can modulate local immune responses in the tumor stroma to support progression and metastasis.

2. Metabolism and tumorigenic properties of PGE₂

PGE₂ makes up the majority of secreted prostaglandin in tumors and is thought to be the principal tumorigenic COX-2-derived product. This has been studied in a broad range of cancers, though perhaps most intensively in colorectal cancer [2]. PGE₂ is generated through the conversion of PGH₂ by microsomal PGE synthases (mPGES) 1 or 2, or cytosolic (c) PGES. Like COX-2, mPGES-1 is inducible and appears to be the dominant PGE₂-generating enzyme in tumors [13]. Functional coupling of COX-2 and mPGES-1 has been reported [14] while the constitutive cPGES couples to COX-1 (mPGES-2 has yet to be well characterized). PGE₂ acts through four distinct G-protein coupled receptors termed EP1, EP2, EP3, and EP4. Regulation of prostaglandin signaling relies not only on their synthesis, but also on their cellular transport and degradation. Solute carrier organic anion transporter 2A1 (SLCO2A1), also known as OAT2A1 or prostaglandin transporter, directs uptake of PGE₂, PGD₂, and PGF_{2α} from the extracellular space into the cytosol. Once there, 15-hydroxyprostaglandin dehydrogenase (15-PGDH) catalyzes the initial step in prostanoid breakdown into their inactive 13,14-dihydro-15-keto-metabolites [15]. The multidrug

resistance protein 4 (MRP4) can transport PGE_2 and $PGF_{2\alpha}$ from the intracellular to the extracellular space [16] and thus may contribute to elevated PGE_2 levels and EP receptor activation. Coordinated regulation of these multiple steps in PGE_2 biosynthesis, metabolism and function, ultimately determines the biological response.

The tumorigenic properties of PGE₂ have been thoroughly reviewed elsewhere [2,4,17], including an in-depth analysis of how PGE2 contributes to the hallmarks of cancer [18,19] self-sufficiency in growth signals, insensitivity to antigrowth signals, evasion of apoptosis, limitless replicative potential, sustained angiogenesis, and tissue invasion/metastasis. Briefly, PGE₂ enhances Wnt signaling through EP2-mediated suppression of glycogen synthase kinase (GSK) 3β [20]. Subsequent accumulation of the β-catenin/T cell factor 4 complex leads to transactivation of perixosome proliferator-activated receptor (PPAR) & and upregulation of pro-oncogenic genes [21]. GSK3β suppression is mediated by activation of phosphoinositide 3-kinase and Akt. In addition, Ga_s which couples to EP2, complexes with Axin, dissociating it from the β -catenin destruction complex, leading to further enhancement of the Wnt signaling pathway [20]. PGE₂ also promotes cell survival by induction of the anti-apoptotic protein Bcl-2 via Ras-MAPK signaling [22], an effect that is partially mediated by PGE₂ transactivation of extracellular growth factor receptor [23]. Multiple studies have implicated PGE₂ production in tumors or tumor cell lines in increased expression of vascular endothelial growth factor and its receptors [24,25], an effect that appears mediated by G_q-coupled EP3 [26,27] signaling through protumorigenic extracellular signaling-regulated kinase/c-Jun N-terminal kinases [28].

3. COX-2 in breast cancer

Animal and human studies report COX-2 overexpression in breast cancer [4,29–31], and strongly support a role for this enzyme in disease progression. Targeted overexpression of COX-2 gene in the mammary epithelium, via the mouse mammary tumor virus, was sufficient to induce mammary tumorigenesis in multiparous mice through a PGE₂-EP2 pathway [32,33]. Further studies in this model revealed an upregulation of cytochrome P450 aromatase that was reversed following COX-2 inhibition with celecoxib [31]. COX-2 inhibition reduced tumorigenesis across a wide range of animal breast cancer models. These have been reviewed extensively [2,4]. Briefly, celecoxib and rofecoxib, both considered selective for COX-2 inhibition, suppressed mammary tumorigenesis in rats treated with 7,12-dimethylbenzanthracene and N-methyl-N-nitrosourea [34,35]. The same inhibitors reduced disease in HER2/neu- and Lewis lung carcinoma (3LL) xenograft-induced models [36,37]. In many studies the molecular mechanism of reduced tumorigenesis has not been defined, other than to note a reduction in PGE₂ signaling on mitogenic and anti-apoptotic pathways. Reduced multiplicity and size of HER2/neu-driven mammary tumors in global COX-2 knock-out (KO) mice was attributed to a concurrent suppression in tumor angiogenesis [38], consistent with the reported contribution of COX-2-derived PGE₂ to the angiogenic switch in mammary tumors that allows disease progression [39].

We have used Cre recombinase technology to target deletion of COX-2 expression selectively to the mammary epithelium. Significantly delayed tumorigenesis was observed independent of modified angiogenesis but coincident with a change in the number and

phenotype of tumor infiltrating cells [40]. These, and other studies [41], indicate a wider role for COX-2 in control of tumor progression via regulation of the microenvironment.

4. Immune regulation of tumorigenesis

In the past decade, evidence has quickly mounted that genetic mutations in classical cancer signaling pathways of tumor epithelial cells cannot fully explain differences in phenotype and clinical development of tumors [42,43]. Indeed, cancer is increasingly considered a disease of the tissue and its progression depends on the supportive or suppressive nature of the microenvironment in the surrounding stroma. The microenvironment contains several different cell types, including fibroblasts, endothelial cells, and immune cells, along with all mediators they release. Cells involved in immune surveillance of tumors include T cells, antigen-presenting cells (APC) such as macrophages and dendritic cells (DC), mast cells, natural killer (NK) cells, myeloid-derived suppressor cells (MDSC), neutrophils, and others [12]. Immune regulation of tumorigenesis and metastasis has been reviewed extensively by others [12,42]. Leukocytic infiltration in tumors was originally considered an attempt by the immune system to reject the tumor. Indeed, this may be the original purpose of cells that have migrated to a tumor site. However, tumors can "hijack" this process by re-educating the immune response, leading to suppression of tumoricidal and immune competent phenotypes, as well as promotion of alternative immune cell functions, often important for normal development and wound healing, that support tumor growth [44,45]. A prominent example of alternative activation is the CD4⁺ T helper lymphocyte type 1 and type 2 (T_H1 and T_H2, respectively) paradigm, in which appropriately stimulated T_H1 cells release cytokines that support cytotoxicity while T_H2 cells contribute to adaptive immunity and humoral antibody production. T_H1-derived cytokines can suppress T_H2 cytokine responses, and vice versa. These complex differential responses by immune cells underscore the need to understand not only the tumor cells themselves, but also the immune responses occurring in the tumor stroma (Fig. 1).

5. Effect of PGE₂ on tumor-associated macrophages

Tumor-associated macrophages (TAM) represent the majority of tumor infiltrating leukocytes [46]. Macrophages are particularly relevant to the tumor microenvironment because, in addition to their role as modulators of angiogenesis, tumor cell migration, and matrix remodeling, they can support or suppress local immune responses thereby contributing to immune surveillance or escape. Macrophages express Toll-like receptors, mannose receptors, and scavenger receptors that can all activate classical innate immunity [47]. They also have adhesion receptors that allow interfacing of macrophages with other immune cells and major histocompatability complexes (MHC) for antigen-presenting functions. In addition, macrophages express a variety of cytokine receptors that can activate signal transducer and activator of transcription (STAT) signaling and induce further release of cytokines [48]. Through their direct and indirect effects on other immune cells, macrophages play a central role in the orchestration of the localized tumor immune response that can support or suppress tumor growth.

As early as the 1970s, macrophages stimulated with lipopolysaccharide (LPS), were shown to kill tumor cells in vitro [49]. These experiments supported the general notion that macrophage activation would be tumor suppressive, a hypothesis reinforced by reports of cytokine-induced release of reactive nitrogen and oxygen intermediates from macrophages [50]. In 1992, Gordon and colleagues coined the term "alternatively activated macrophages" after noting that stimulating macrophages with interleukin (IL)-4 inhibited pro-inflammatory cytokine production and restricted expression of MHC class II complexes [51]. These two unique polarization states have since been termed M1 (classically activated) and M2 (alternatively activated), so-named after the T_H1/T_H2 paradigm, and seem to represent two poles of macrophage function. Indeed, research within the past decade has revealed that M2 macrophages, which share many features of TAMs [52], support cell growth, tissue remodeling, and angiogenesis, and are considered to be pro-tumorigenic [53]. In addition, whereas M1 macrophages release type 1 cytokines, such as tumor necrosis factor (TNF) a and IL-6, that promote T lymphocyte differentiation to pro-immune CD4⁺ T_H1 and cytotoxic T lymphocytes (CTL; CD8⁺ T lymphocytes), M2 macrophages release immunosuppressive cytokines such as IL-10 and TGFβ that result in T cell anergy. M1 and M2 macrophages also utilize distinct arginine-metabolizing enzymes that characterize their polarization state and contribute to their distinct functions; M1 macrophages express inducible nitric oxide synthase (iNOS), metabolizing L-arginine into nitric oxide thus contributing to M1-associated tumoricidal functions. M2 macrophages, in contrast, express arginase-1, which metabolizes L-arginine into L-ornithine and proline, benefiting cell growth and promoting extracellular matrix remodeling. Further, the markedly enhanced expression of arginase-I in M2 macrophages depletes the pool of local L-arginine that is necessary for normal CTL function [54].

PGE₂ is typically associated with immunosuppression, restraint of type 1 cytokine production and M1 macrophage polarization, as well as enhanced expression of M2 markers. In early work, treatment of Lewis lung carcinoma (LLC) tumor bearing mice with indomethacin, a mixed COX-1/COX-2 inhibitor, attenuated tumor suppressive natural killer (NK) cells, an effect that was associated with macrophages and PGE₂ [55]. Additional studies support a role for PGE₂ in the control of macrophage phenotype and function. A recent report that COX-2 inhibition reversed tumor-mediated bone marrow-derived cell differentiation to APCs revealed an autocrine function of COX-2 derived mediators in regulation of myeloid cell phenotype [48]. Interestingly, suppression of the APC phenotype by tumor-derived factors was coincident with a decrease in bone-marrow cell expression of 15-PGDH/SLCO2A1 mRNA and an increase in MRP4 mRNA, suggesting a coordinated effort at multiple points in the COX-2-PGE₂ metabolic pathway. However, inhibition of mPGES-1, the dominant source of macrophage PGE₂ [56], did not recapitulate the phenotypic rescue observed with a COX-2 inhibitor, suggesting significant complexity in the biology, perhaps involving other COX-2-derived prostanoids [48]. STAT1/3 signaling in macrophages, a known cytokine regulatory pathway, was implicated in COX-2-mediated suppression of APC function. Suppressed function of other inflammatory/immune signaling pathways has been linked with COX-2 and PGE2. PGE2 can induce expression and homodimerization of the inhibitory p50 subunit of nuclear factor (NF)-κB, resulting in decreased NFxB activation and depressed M1 cytokine release in TAMs isolated from

human ovarian cancers [57]. PGE_2 also decreased elaboration of several type 1 cytokines, including IL-1 β [58], TNF α , and IL-6 [59] in LPS-stimulated human peripheral blood monocytes via EP2/EP4-cAMP signaling, IL-8 in LPS-stimulated human alveolar macrophages [60], and IL-6 and TNF α in LPS-induced murine peritoneal macrophages, again through an EP2 and EP4 cAMP-mediated pathway [61]. PGE_2 also decreased expression of iNOS after LPS stimulation in J774 cells [62].

Conversely, PGE $_2$ is implicated in the promotion of M2 macrophage function, promoting expression of the M2 cytokine IL-10 and arginase-1 in LPS-stimulated murine peritoneal macrophages [61] and bone marrow-derived macrophages [63]. Enhanced PGE $_2$ metabolism, via forced 15-PGDH overexpression, reduced secretion of IL-10, IL-13, and IL-6, with a coincident increase in F4/80⁺/CD11c⁺ APC and a decrease in F4/80⁺/CD11c⁻M2 macrophages in a xenograft model of colon cancer. Experiments in our lab examining the role of PGE $_2$ in modulating M1 and M2 polarization reinforce these findings.

Together, these data support autocrine and paracrine effects of PGE₂, and possibly other COX-2-derived prostanoids, in the restraint of M1, and/or promotion of M2, macrophage function. The studies outlined above describing COX-2 and PGE₂ modulation of macrophage and APC functions reinforce the paradigm that PGE₂ downregulates expression of M1 markers/cytokines (e.g. TNFα, IL-6, and iNOS) and functions (e.g. antigen-presenting ability and release of NO), while upregulating M2 markers/cytokines (IL-10 and arginase-1). Much of this work has been in *in vitro* or *ex vivo* models. Additional research is warranted to determine the contribution of COX-2 and PGE₂ to regulation of M1/M2 polarization *in vivo* across the entire spectrum of tumor onset and progression. Indeed, while TAMs have many of the characteristics of M2 macrophages, they are a distinct population. In particular, the high expression levels of COX-2 in TAM diverges from the typical M2 phenotype in which COX-2 is suppressed, underscoring the complex influence of tumorassociated mediators and COX-2-derived prostanoids in paracrine and autocrine control of macrophages.

6. Effect of PGE₂ on myeloid-derived suppressor cells

A growing appreciation for the role of MDSCs in tumor immunosuppression has developed over the past decade. Interest in MDSCs began in the 1980s when MDSCs were more commonly referred to as natural suppressor cells or bone marrow suppressor cells. MDSCs were originally identified as a subset of bone marrow cells that could inhibit T cell and NK proliferation that were characteristically distinguished from macrophages by their inability to adhere to nylon wool [64]. Since then, murine MDSCs have been defined as CD11b⁺, Gr-1⁺, F4/80⁺, CD11c^{low} cells that express both arginase-1 and iNOS [65]. MDSCs include a variety of myeloid progenitor cells as well as modified myeloid-derived cells, including immature macrophages, DCs, and granulocytes (such as neutrophils).

MDSC regulation of the immune system, especially as it pertains to cancer, has been well reviewed [65]. In short, cytokine receptor activation can signal through the Janus kinase (JAK)-STAT pathway to regulate cell survival and activation. STAT3 is particularly important in tumor-dependent expansion of MDSCs, as STAT3 regulation of gene

transcription can lead to Myc, Bcl-XL, and cyclin D1 transcription, enhancing MDSC survival. Leukocyte-derived type 1 and 2 cytokines signal primarily through STAT1 and STAT6, respectively, enhancing the immunosuppressive environment through upregulation of TGFβ, IL-10, arginase-1, and iNOS in MDSCs. Dual expression of both arginase-1 and iNOS differentiates MDSCs from other mature myeloid cells. Both of these enzymes use L-arginine as a substrate, and L-arginine depletion inhibits T cell proliferation through a variety of mechanisms [54]. In addition, NO production inhibits JAK–STAT signaling in T lymphocytes, inhibits antigen-presenting functions of MHC II expressing cells, including macrophages [66,67], and causes T cell apoptosis [68,69]. Importantly, reactive oxygen species (ROS) production through arginase-1 and iNOS can produce peroxynitrite, which is directly linked to insensitivity of CD8⁺ CTLs to antigen [70]. Given their multiple and potent immunosuppressive effects, MDSCs may provide a valuable target in host immune defense against tumorigenesis.

PGE₂ has been strongly implicated in MDSC function with respect to chemoattraction of MDSCs to tumors sites as well as differentiation of non-bone marrow cells to MDSCs in the tumor microenvironment. MDSCs positive for both CD11b and Gr-1 derived from 4T1 mouse mammary tumors expressed all four EP receptors [71]. In addition, treatment of bone marrow cells with PGE₂ or EP2 selective agonist (butaprost) increased differentiation to MDSC by 20% and 40%, respectively, while antagonism of EP1/EP2 or EP4 prevented differentiation into MDSC [71]. PGE₂-induced elevation in MDSC was coincident with decreased T cell activation [71]. In addition, reduced 4T1-xenograft tumor growth in EP2^{-/-} mice was coincident with decreased MDSC infiltration in tumors [71]. Another 4T1 injection model replicated these results, showing that PGE₂ and TGF β were both necessary for MDSC differentiation from bone marrow cells [72]. Further, PGE₂ concentrations in tumor cell supernatants correlated very strongly with ability to induce MDSC differentiation in BM cells and MDSC differentiation was offset when PGE₂-depleting antibody was added to the culture mixture [72].

In 3LL injection models of murine tumorigenesis, C57BL/6 MDSCs expressed similar levels of arginase-1 as those from severe combined immunodeficiency mice, indicating that arginase-1 expression may not be dependent on T cell-derived factors [73]. Instead, MDSCs cultured with tumor cells or conditioned medium showed a significant increase in arginase-1 expression. PGE₂ was identified as a dominant soluble tumor-derived mediator driving arginase-1 expression via an EP4/cAMP signaling cascade [73]. In another 3LL tumor cell study, PGE₂ released downstream of Fas ligation increased chemoattraction of MDSCs implicating COX-2 and PGE₂ in Fas-dependent MDSC tumor infiltration [74]. COX-2 inhibition with celecoxib prolonged survival of 1,2 dimethylhydrazine-injected mice coincident with a decrease in Gr-1⁺, CD11b⁺ MDSCs and a decrease in iNOS and arginase-1 mRNA [75]. Celecoxib treatment in mesothelioma-injected mice also showed a decreased expansion of MDSCs coincident with decreased PGE₂ concentrations and improved DC-based co-therapy [76]. Thus, the chemopreventative benefits associated with COX-2 inhibition likely involves repressed MDSC differentiation, infiltration, and/or function.

Intensive research on PGE₂ dependent regulation of MDSC function tumors continues with a strong consensus building that PGE₂ supports MDSC proliferation and consequent immunosuppression. While findings *in vitro* and in tumor xenograft models are promising, further *in vivo* modeling of PGE₂ and MDSC interactions will determine the viability of MDSC-related targets in cancer prevention or treatment via the COX-2 pathways.

7. Effect of PGE₂ on T regulatory cells

A subset of CD4⁺ T cells that suppress effector T cell functions have a crucial role in preventing self (vs. non-self) destruction and thus are implicated in a variety of autoimmune diseases [77]. This same subset of T cells may be involved in T cell anergy in tumorigenesis and modify the immune response to accept the tumor as self instead of marking it for destruction. Classification, development, and function of these regulatory T cells (T_{ress}), which is still ill-defined and the subject of much research, has been reviewed elsewhere [77,78]. CD4+CD25+FOXP3+ cells are generally accepted as a population of T_{reg}, and strength of interaction with T_{reg} MHC complexes appear crucial to T_{reg} development. T_{reg} suppressor function has been expanded to encompass many of the effector cells of the immune system, including T_H cell subsets, CD8⁺ CTLs, macrophages, NK cells, and mast cells [79]. The exact mechanism of T_{reg} suppression is still under intensive investigation although IL-10, TGFβ, and IL-35 suppressive cytokines are considered characteristic of T_{regs}, as are interactions between lymphocyte-activating gene 3 and MHC II complexes in DCs leading to suppressed DC maturation and limited APC functions [78]. Potential cancer immunotherapies targeted at T_{regs} include TGF β depletion and stimulation of glucocorticoid-induced TNF receptor to induce T effector cell differentiation [79].

In gastric cancer patients, the proportion of CD4+CD25+FOXP3+ cells was increased in both the peripheral blood and tumor tissues as compared to normal healthy patients [80]. This correlated with stage of tumor progression, as well as concentration of PGE2, in ascites of gastric cancer patients but interestingly not with levels of TGF\$\beta\$ and IL-10, both of which are T_{reg} factors. COX-2 expression also strongly correlated with FOXP3 expression by flow cytometry, implicating T_{regs} as PGE₂-producing cells in the tumor microenvironment. FOXP3 protein levels were reduced using both a non-selective COX and COX-2 selective inhibitors, but not using EP2 or EP4 antagonists [80]. However, these antagonists did match COX-2 inhibitors in reversing suppressed proliferation of tumor-infiltrating T lymphocytes. These may indicate that PGE₂ acting on receptors EP2 and EP4 are only partially responsible for T_{reg} differentiation and infiltration and/or other COX-2-derived factors contribute to effector T cell suppression. In a separate study using CD4⁺CD25⁺ T_{regs} isolated from human peripheral blood, stimulation with anti-CD3/CD28 led to a large induction of PGE₂, IL-10, and TGFβ. Addition of COX-2 inhibitor, anti-IL-10, or anti-TGFβ reversed T_{reg} suppression of IFNγ-producing CD4⁺ T cells, a measure of effector T cell function. Concordant with the correlation COX-2 and FOXP3 expression [80,81], treatment with PGE2 dose-dependently increased FOXP3 expression in peripheral blood T_{regs}, as well as in colon cancer patients [82]. These studies emphasize a role for PGE₂ mediating T cell proliferation in both an autocrine $T_{reg} \rightarrow T_{reg}$ and paracrine $T_{reg} \rightarrow$ effector T cell manner.

In an injection model of non-small cell lung cancer, knockdown of COX-2 in tumor cells led to decreased T_{regs} at the tumor site, an effect that was mirrored with anti-PGE $_2$ antibody or a selective COX-2 inhibitor [83]. In *ex vivo* studies, COX-2 inhibition increased effector T cell proliferation in co-cultures with murine T_{regs} . Robust upregulation of FOXP3 was seen when CD4+CD25+ cells were pulsed with PGE $_2$, an effect that was partially reduced through the use of EP4-/- cells and completely ablated in EP2-/- cells [83]. These studies provide further evidence that COX-2-derived PGE $_2$, acting through EP2 and EP4, increases T_{reg} infiltration, maturation, and function, leading to immunosuppression and increased tumor cell survival.

In general, there is a consensus that PGE_2 can enhance T_{reg} proliferation and FOXP3 overexpression with consequent suppression of effector T cell proliferation. However, reports that PGE_2 can instead suppress proliferation of T_{regs} [84,85] reveal the complexity of COX-2 and PGE_2 in regulation of T_{reg} differentiation, proliferation, and contribution to disease in the already immunosuppressive environment of a growing tumor.

8. Conclusion

The tumor immune microenvironment involves a remarkably complex interplay of cells and mediators that can drive a tumor towards limitless growth or imminent destruction. Multiple different cell types, and subtypes, exist in a milieu of growth factors, cytokines, oxidative species, and lipid mediators. Among these, PGE₂ has emerged as a mediator that not only impacts classical oncogenic signaling pathways in tumor cells, but also contributes to shifting the tumor microenvironment towards immune suppression and evasion, promoting tumorigenesis. A substantial body of work indicates the immune suppressive role of PGE₂, including the restraint of M1, and promotion of M2, macrophage phenotypes, augmentation of MDSC differentiation and infiltration, and support of T_{reg} proliferation and function. These immunosuppressive leukocyte subtypes share common cytokine mediators, including TGFβ and IL-10. More importantly, each cell subtype can generate PGE₂, providing an autocrine mechanism for prolonging and enhancing their own immunosuppressive phenotype. This emphasizes the importance of exploring the tumor microenvironment as a whole, rather than focusing on alterations in an individual subset of tumor-associated cells. There are several other types of immune cells that have not been discussed above, including B cells, mast cells, eosinophils, and neutrophils, which also appear to contribute to immune surveillance of tumorigenesis. Indeed, an N1/N2 tumor-associated neutrophil paradigm has recently been discussed [86]. Discrepancies between the immunosuppressive actions of PGE₂ across in vitro or ex vivo experiments underscore the context-specific nature of the tumor microenvironment, a context that is unique to individual cancers and disease stages. Continuing research in COX-2-derived PGE₂-based immunosuppression may yield appealing targeted immunotherapies to reclaim the immune system's ability to destroy a cancer.

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Abbreviations

COX cyclooxygenase
PG prostaglandin

mPGES microsomal prostaglandin E synthase

15-PGDH 15-hydroxyprostaglandin dehydrogenase

SLCO2A1 solute carrier organic anion transporter 2A1

APC antigen-presenting cell

DC dendritic cell
NK natural killer

MDSC myeloid-derived suppressor cell

TAM tumor-associated macrophage

CTL cytotoxic T lymphocyte

iNOS inducible nitric oxide synthase

 T_{regs} regulatory T cells

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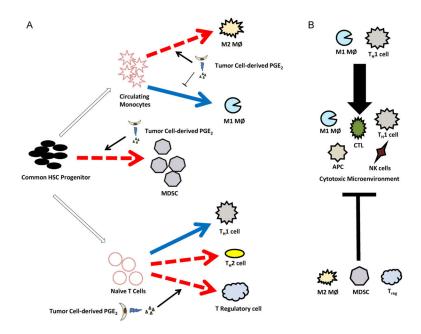


Fig. 1. Schematic of how PGE_2 may affect immune cell differentiation and function in tumors. (A) Immune cells of the microenvironment may be immunogenic (solid arrows) or immunosuppressive (dashed arrows). Tumor cell-derived PGE_2 can affect the differentiation, infiltration, and maturation of several tumor-associated immune cells. PGE_2 suppressed polarization of macrophages (MØ) to the M1 phenotype while enhancing M2 characteristics. PGE_2 promotes differentiation to and function of MDSCs and is correlated with increased MDSC numbers. PGE_2 has been implicated in increased differentiation and infiltration T_{reg} which in turn suppress the maturation of other T cells. (B) These actions of PGE_2 work to shift the tumor microenvironment into an immunosuppressive environment by suppressing maturation of cytotoxic cells that are essential for immune surveillance.