

Review Article

The Signaling Pathway of PGE₂ and Its Regulatory Role in T Cell Differentiation

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Prostaglandin E₂ (PGE₂) is a lipid mediator derived from the fatty acid arachidonic acid. As an essential inflammatory factor, PGE₂ has a critical impact on immune regulation through the prostanoid E (EP) receptor pathway. T cells, including CD4⁺ and CD8⁺ T cell subsets, play crucial roles in the adaptive immune response. Previous studies have shown that PGE₂ is involved in regulating CD4⁺ T cell differentiation and inflammatory cytokine production via the EP receptor pathway, thereby affecting the development of diseases mediated by CD4⁺ T cells. In this review, we summarize the signaling pathway of PGE₂ and describe the relationship between PGE₂ and T cell differentiation. Hence, this review may provide important evidence for immune therapies and may even promote the development of biomedicines.

1. Introduction

Prostaglandin (PG) is a lipid mediator family derived from the fatty acid arachidonic acid (AA). Due to differences in their molecular structures, PGs were classified into nine groups, including PGA, PGB, PGD, PGE, PGF, PGG, PGH, PGI, and PGJ. Among these molecules, PGE₂ is a kind of inflammatory factor that has been intensively studied. Although PGE₂ has a short half-life and a 90% degradation rate in pulmonary circulation [1], it plays a significant role in mediating inflammation and multiple physiological processes [2]. PGE₂ functions by prostanoid E (EP) receptors, which include four types of membrane-bound G protein-coupled receptors named EP1 to EP4 mediating the multiple signaling pathway. To date, a series of studies has demonstrated that PGE₂ regulates the differentiation, maturation, and activation of immune cells, especially T cells [3, 4].

As essential components in the adaptive immune system, T cells are mainly divided into CD4⁺ helper T (Th) cells and CD8⁺ cytotoxic T lymphocytes (CTLs) [5]. After stimulation, CD4⁺ T cells can differentiate into a variety of effector subsets, including classical Th1 cells and Th2 cells, the subsequently defined Th17 cells, follicular helper T (Tfh)

cells, and induced regulatory T (iTreg) cells. Th1 cells are characterized by their secretion of IFN- γ and are involved in cellular immunity, while Th2 cells produce IL-4, IL-5, IL-10, and IL-13 and are required for humoral immunity and allergic reactions. Th17 cells mainly secrete IL-17A, IL-17F, IL-21, and IL-22 and mediate inflammatory response. Tfh cells are a new subset of helper T cells that produce IL-21 and regulate the maturation of B cell responses. Treg cells are characterized by the expression of the forkhead transcription factor (Foxp3) and have essential roles in the maintenance of immune homeostasis by suppressing these effector T cell responses [6]. CTLs play their roles by directly killing infected cells and cancer cells. In this review, we provide a general overview of the metabolism of PGE₂, the signaling pathways of EPs, and the regulatory role of PGE₂ in T cell differentiation.

2. PGE₂ Synthesis and Metabolism

Prostaglandins are derived from AA and synthesized mainly through the cyclooxygenase (COX) pathway (Figure 1). When cells are damaged or stimulated, AA is released from plasma membrane phospholipids [7]. Free AA can be

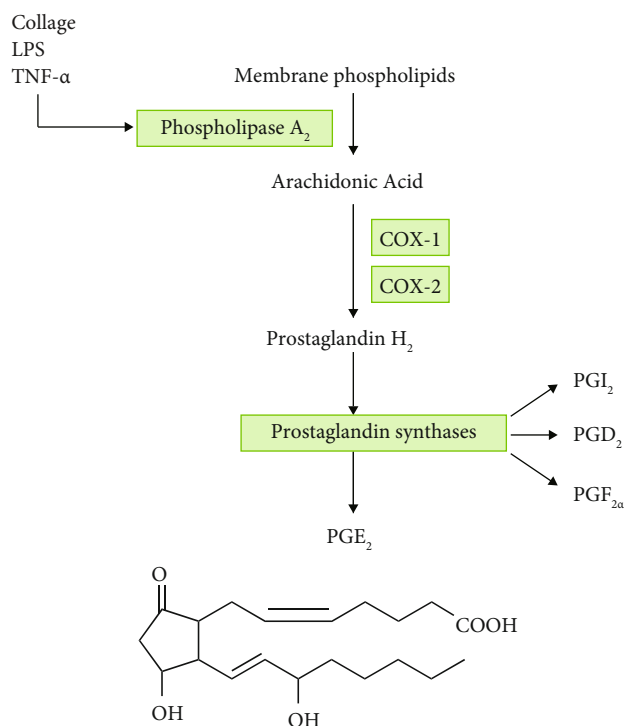


FIGURE 1: The process of PGE₂ synthesis. Arachidonic acid is released from the membrane by phospholipase A₂. Cyclooxygenases (COX-1 and COX-2) produce prostaglandin H₂ (PGH₂) from arachidonic acid. PGH₂ is produced by prostaglandin synthase to produce PGI₂, PGD₂, PGE₂, and PGF_{2α}.

modified into prostaglandin or leukotrienes [8]. For prostaglandin synthesis, AA released from membranes is oxidized by the cyclooxygenase enzymes COX-1 and COX-2 [9, 10]. The enzymatic effect of COX enzymes on AA involves two steps, one is the oxidation of AA to form prostaglandin G₂ (PGG₂), and the other is the reduction of PGG₂ to form prostaglandin H₂ (PGH₂). Depending on the specific enzymes, PGH₂ can be modified to five different prostaglandins, including PGD₂, PGI₂, PGF_{2α}, PGE₂, and thromboxane A₂ (TXA₂) [2, 7]. PGE₂ synthesis through COX is mediated by three distinct enzymes, cytosolic PGE synthase (cPGES), microsomal PGE synthase-1 (mPGES-1), and mPGES-2 [2].

Nearly all types of cells and tissues can secrete PGs. When PGs are secreted, however, they are quickly degraded by the lung or liver. The lung is the primary organ involved in PGE₂ metabolism [11, 12]. PGE₂ metabolism mainly occurs via three mechanisms, leading to the inactive metabolite: (i) 15-hydroxyprostaglandin dehydrogenase (15-PGDH) converts a 15-hydroxyl group to a keto group for metabolism; (ii) 9-ketoreductase reduces 9-ketone to hydroxyl; and (iii) a side chain is inactivated by β -oxidation and/or ω -hydroxylation. The final metabolites of PGE₂ are eliminated from the body in the urine. The effects of PGE₂ are regulated by the balance between its COX-2-regulated synthesis and 15-PGDH-driven degradation [13].

2.1. PGE₂ Signaling and EP Receptors. After PGE₂ is synthesized in the cytoplasm, it diffuses out of the cell through

facilitated diffusion. Since PGE₂ is quickly degraded, it is distinct from typical hormones that act on distant target tissues but can only be produced and released locally. After being secreted from cells, the lipid mediator binds to membrane receptors, activates downstream signaling pathways, and exerts biological effects [14].

PGE₂ signals through four morphologically distinct G-protein-coupled receptors, namely, EP1, EP2, EP3, and EP4, and triggers divergent signaling pathways (Figure 2), thereby mediating its various biological functions. The mRNAs of the EP receptors also exhibit different expression patterns in a number of tissues, and activating each receptor subtype leads to distinct functional consequences [15]. EP3 and EP4 are high-affinity receptors and are expressed in all human tissues, whereas the activation of EP1 and EP2 occurs in only a few organs and requires significantly higher levels of PGE₂ [16]. EP receptors are also expressed on various immune cell membranes. EP2 and EP4 are the main receptors of PGE₂ involved in regulating the differentiation of CD4⁺ T cells [17]. By coupling with the activated G protein, it stimulates adenylate cyclase (AC) to activate the cAMP/protein kinase A (PKA)/cAMP-responsive element-binding protein (CREB) signaling pathway and downstream molecules to participate in mediating proinflammatory responses and inhibiting PGE₂ activity [18]. It is well known that clear signal transduction of EPs is beneficial for therapeutic strategies against diseases related to the immune system.

2.2. EP1 Receptor. In humans, the EP1 receptor has the lowest affinity for PGE₂ [19] and couples to a Gq alpha subunit (G α q). PGE₂ can result in smooth muscle contraction through this complex. The human single EP1 receptor consists of 402 amino acid residues, whereas the rat single EP1 receptor consists of 405 amino acid residues. The EP1 receptor has 7 hydrophilic transmembrane domains. An arginine residue exists in the 7th domain, and this residue is an important PGE binding site. The phosphorylation of PGE₂ depends on cyclic adenosine monophosphate- (cAMP-) dependent protein kinase and protein kinase C (PKC). The G α q subunit activates phosphoinositide-phospholipase C (PLC) and finally leads to an elevation in intracellular Ca²⁺ and the activation of PKC, mediating gene transcription through the activation of the nuclear factor of activated T cells (NFAT), nuclear factor-kappaB (NF- κ B), and the mitogen-activated protein kinase (MAPK) pathways [16].

EP1 receptor mRNA is highly expressed in the kidney, lung, and stomach [15]. EP1 has not been demonstrated to play a significant role in innate immunity but participates in controlling T cell differentiation [2, 4].

2.3. EP2 Receptor. The human EP2 receptor consists of 358 amino acids [20]. This receptor is coupled to a Gs alpha subunit (G α s). The long C-terminal tail of the EP1 receptor contains serine and threonine, the phosphorylation of which depends on cAMP-dependent protein kinase [15, 21]. Gs proteins lead to an increase in cAMP and activation of PKA. A high concentration of intracellular cAMP can also activate both PKA and the exchange protein directly activated by cAMP (EPAC). Then, EPAC phosphorylates the

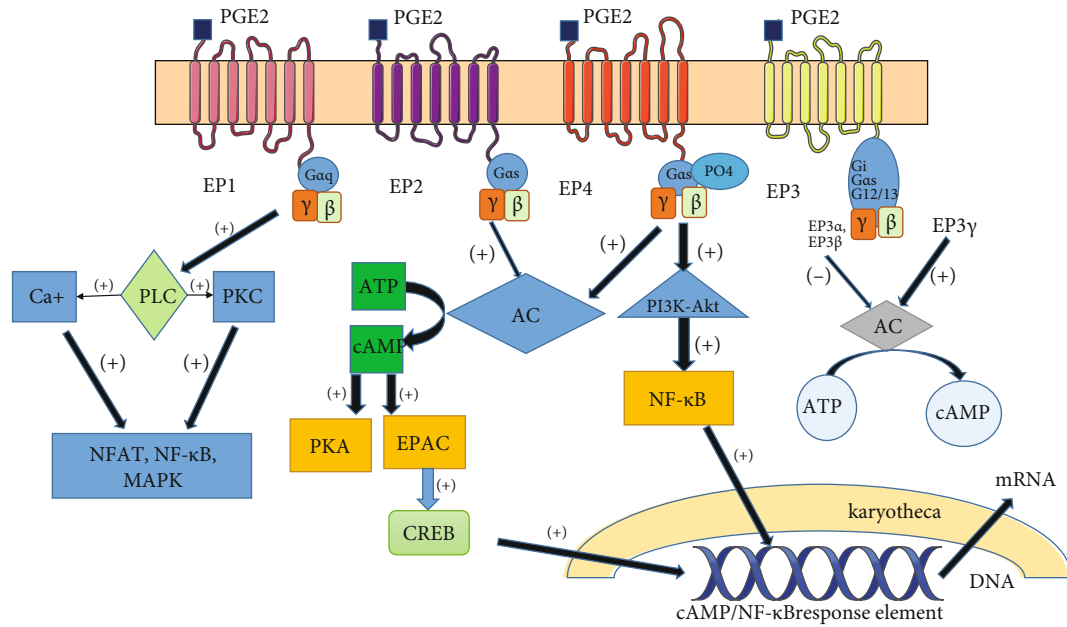


FIGURE 2: The signaling pathway of PGE₂. The schematic diagram shows the downstream signaling pathway of PGE₂. The lipid mediator activates or inhibits the target cell via four EP receptors, EP1, EP2, EP3, and EP4. Different receptors activate different signaling pathways which are connected in a network in the cell and cooperate with each other in vital processes.

transcription factor CREB [22]. In addition, EP2 receptor signaling leads to the inhibition of glycogen synthase kinase-3 (GSK-3), which inhibits the translocation of β -catenin into the nucleus [21].

The EP2 receptor participates in most of the immunoregulatory effects of PGE₂ in both the innate and adaptive immune responses. For example, PGE₂ in the supernatant of thyroid cancer cells inhibits the activity of NK cells through the EP2 and EP4 receptors [23]. In addition, PGE₂ promotes neurite outgrowth and suppresses cell proliferation by activating the EP2 receptor subtype, and the cAMP signaling pathway is involved in the PGE₂-induced differentiation of NSC-34 cells [24].

2.4. EP3 Receptor. The EP3 receptor, which consists of 365 amino acid residues, is an abundantly and widely expressed EP receptor [15]. EP3 also contains two cAMP-dependent protein kinase phosphorylation sites. EP3 is the only EP receptor that includes multiple variants (α , β , and γ). During the transcription of the EP3 coding sequence, several types of mRNA spliceosomes are produced. The differentiation of these spliceosomes results in a change in the C-terminal tail [25]. It should be noted that these variants have a similar affinity for PGE₂, but they have different signal transduction pathways, relative expression patterns, and desensitization properties [2]. EP3 α and EP3 β are coupled to Gi and inhibit adenylyl cyclase (AC) or cAMP, whereas EP3 γ is coupled to Gs or Gi and stimulates cAMP production. Therefore, a high concentration of PGE₂ inhibits AC, while a low concentration of PGE₂ activates AC [26].

EP3 mRNA can be detected in almost all tissues. EP3 signaling participates in many metabolic processes. The selective inhibition of EP3 might be a potential approach

for reducing chronic neuropathic pain [27]. Moreover, it has been reported that EP3 signaling is induced in placentas associated with unexplained recurrent pregnancy losses [28].

2.5. EP4 Receptor. The EP4 receptor signals through a pathway similar to that of the EP2 receptor. The EP4 receptor consists of 513 amino acids. Similar to EP2, EP4 couples to the Gas protein and activates the cAMP/PKA pathway. cAMP is rapidly degraded by phosphodiesterase to limit the duration of the signal. PKA then phosphorylates target proteins in the cells [29]. EP2 exhibits higher sensitivity and can more effectively increase cAMP. However, the EP4 receptor has a higher affinity for PGE₂ than the EP2 receptor [21]. Compared to EP2, EP4 performs the remarkable function of activating the phosphatidylinositol 3 kinase (PI3K) signaling pathways. The subsequent phosphorylation is mediated via G-coupled receptor kinases or via the ability to bind Gi protein [30, 31]. In addition, EP4 also stimulates noncanonical activation of the PI3K-Akt (also known as protein kinase B) and extracellular regulated kinase (ERK) pathways [16, 29].

Among all four types of EP receptors, the maintenance of EP4 has received much attention [32]. PGE₂-EP4 interaction causes muscle-specific stem cell expansion by triggering a cAMP/pCREB pathway that activates the proliferation-inducing transcription factor *Nurr1* [33]. Besides, activation of EP4 may mediate many cellular responses, such as the promotion of angiogenesis, proliferation, motility, and metastasis or the delay of tumor cell apoptosis [32].

2.6. Effects of PGE₂ on T Cell Subsets. PGE₂ exerts many complex immunoregulatory roles under physiological and pathophysiological conditions [34, 35]. PGE₂ influences the

differentiation of effector T cells, such as Th1, Th2, Th17, Treg, Tfh, and CTLs [3] (Figure 3).

2.7. PGE₂ and Th1 Cells. Th1 cells mainly secrete IFN- γ , IL-2, and TNF- α , mediating cellular immune responses and playing a key regulatory role in resistance to bacterial and viral infections. T-bet is the key transcription factor of this T cell subset. Previous experiments indicated that both EP2- and EP4-mediated Th1 differentiation are inhibited by PI3K inhibitors [36]. This signaling pathway can decrease the cAMP level and then reduce the cAMP-mediated inhibition of T cell, indicating that PGE₂ could activate the differentiation of Th1 cells. Other researches suggested that PGE₂ selectively inhibits the differentiation of naïve CD4⁺ T cells into Th1 cells and noted that PGE₂ can also decrease the production of IL-12 by monocytes or dendritic cells, leading to an effect on the proliferation and differentiation of Th1 cells [37, 38].

2.8. PGE₂ and Th2 Cells. Th2 cells, which mediate humoral immunity, mainly secrete IL-4, IL-5, IL-10, and IL-13. GATA-3 is considered the crucial transcription factor of Th2 cells, autoactivating its expression and driving epigenetic changes in the Th2 cytokine cluster (*IL4*, *IL5*, and *IL13* genes) while suppressing the factors critical for regulating the Th1 pathway, such as signal transducer and activator of transcription factor 4 (STAT4) and the IL-12R β 2 chain [39]. PGE₂ inhibits the production of both IL-4 and IL-5 by Th2 clones [40]. Later, Bao et al. discovered that PGE₂ can enhance the IL-4/IFN- γ ratio in CD4⁺ T cell culture and polarize CD4⁺ T cells towards Th2 cells [41]. Hence, the promotion effect of PGE₂ mainly activates the differentiation of Th2 cells via the inhibition of Th1 cells.

2.9. PGE₂ and Th17 Cells. Th17 cell differentiation requires ROR γ t, a transcription factor that is induced by TGF- β in combination with the proinflammatory cytokines IL-6, IL-21, and IL-23, all of which activate STAT3 phosphorylation. The differentiation of naïve CD4⁺ T cells into Th17 cells is mediated by the EP2 or/and EP4 receptor leading to cAMP production. Moreover, PGE₂ increases the concentrations of IL-23R and IL-1 β R. cAMP and other cytokines induce the expression of IL-23R via the EP2 receptor [17, 42]. Klasen et al. found that the expression level of EP4 is significantly increased in Th17 cells from patients with ankylosing spondylitis (AS) compared to those in healthy individuals or rheumatoid arthritis (RA) patients and that the EP4 expression level in Th17 cells from AS patients has a positive correlation with disease activity [43]. Lee et al. declared that the T cell-intrinsic EP2/EP4 signaling is critical in IL-23-driven generation of pathogenic Th17 cells and consequent pathogenesis in psoriasis [44].

2.10. PGE₂ and Tfh Cells. Tfh cells are defined as CD4⁺ T helper cells that express CD40L, chemokine receptor 5 (CXCR5), programmed death 1 (PD-1), and inducible T cell costimulator (ICOS). Bcl-6 is the characteristic transcription factor of Tfh cells [45]. Tfh cells migrate into follicles and interact with antigen-specific B cells to support their differentiation into memory B cells or plasma cells [46]. Tfh cells can secrete IL-21, which binds to IL-21R on the surface of B

cells, assists B cell activation and proliferation, and induces self-antibody production. IL-21 is a potent differentiation factor for B cells. Tfh cell defects or hyperactivity can cause immune system dysfunction, leading to the occurrence of autoimmune diseases [47]. A recent study has shown that PGE₂ can facilitate antibody class switching in B cells by inducing Tfh cell differentiation [48]. We recently discovered that the serum concentration of prostaglandin E metabolite (PGEM) and the proportion of Tfh cells are increased in the collagen-induced arthritis mouse model compared with those in wild-type mice. In addition, there is a positive correlation between the concentration of serum PGEM and the population of Tfh cells in RA patients, suggesting that PGE₂ acts as a positive regulator of the process of Tfh differentiation [1]. However, the mechanism underlying the regulation of Tfh differentiation by PGE₂ still needs to be explored.

2.11. PGE₂ and Treg Cells. Tregs are immunosuppressive cells that mainly participate in immune homeostasis by acting as a major barrier to effective immunity against tumors and sterilizing immunity against chronic viral infections [35]. The membrane-bound IL-2 α -chain is a marker of Treg cells. There are two critical cytokines, namely, IL-2 and TGF- β , that drive the differentiation of Treg cells from naïve T cells. IL-2R is required for both the thymic and peripheral generation of Treg cells. TGF- β inhibits the recruitment of Dnmt1, the key maintenance DNA methyltransferase whose activity likely leads to silencing of the newly induced Foxp3 gene [49]. Foxp3 is a specific transcription factor that regulates the development of Treg cells and is essential for the immunosuppressive function of Treg cells, which is initially considered a specific marker of Treg cells. It has been reported that PGE₂ promotes the growth of Treg cells in both humans and mice via the EP2 or EP4 receptor [50]. Since PGE₂ can effectively suppress the EP/cAMP/PKA pathway, it also inhibits the differentiation of Treg cells. It is worth mentioning that PGE₂ signaling via EP2 expressed on human naïve CD4⁺ T cells suppresses Treg differentiation *in vitro* via the cAMP-PKA signaling pathway [3]. However, when UV was used to induce an immunosuppressive reaction, PGE₂ facilitated an increase in Treg cell numbers. Additionally, Kitipong et al. showed the impairment of the immunosuppressive effect of UV with an EP4 antagonist and the reversal of the indomethacin-induced impairment of immunosuppression by an EP4 agonist. Notably, treatment with an EP4 agonist alone, without UV irradiation, does not result in immunosuppression [50]. Additionally, PGE₂ inhibits the expression and production of IL-27 by activating conventional dendritic cells *in vivo* and *in vitro*, thus inhibiting the IL-27-induced differentiation and IL-10 production of murine CD4⁺CD49b⁺LAG-3⁺Foxp3⁺Tr1 cells [51]. The imbalance of Th17/Treg may play a role in the progression of autoimmune diseases such as RA [52]; therefore, PGE₂ is expected to become one of the therapeutic directions.

2.12. PGE₂ and CD8⁺ CTLs. CTLs, which originate from naïve CD8⁺ T cells, can specifically recognize endogenous antigen peptides (i.e., MHC I complex) and induce the apoptosis of target cells. Li et al. showed that CD8⁺ T cell

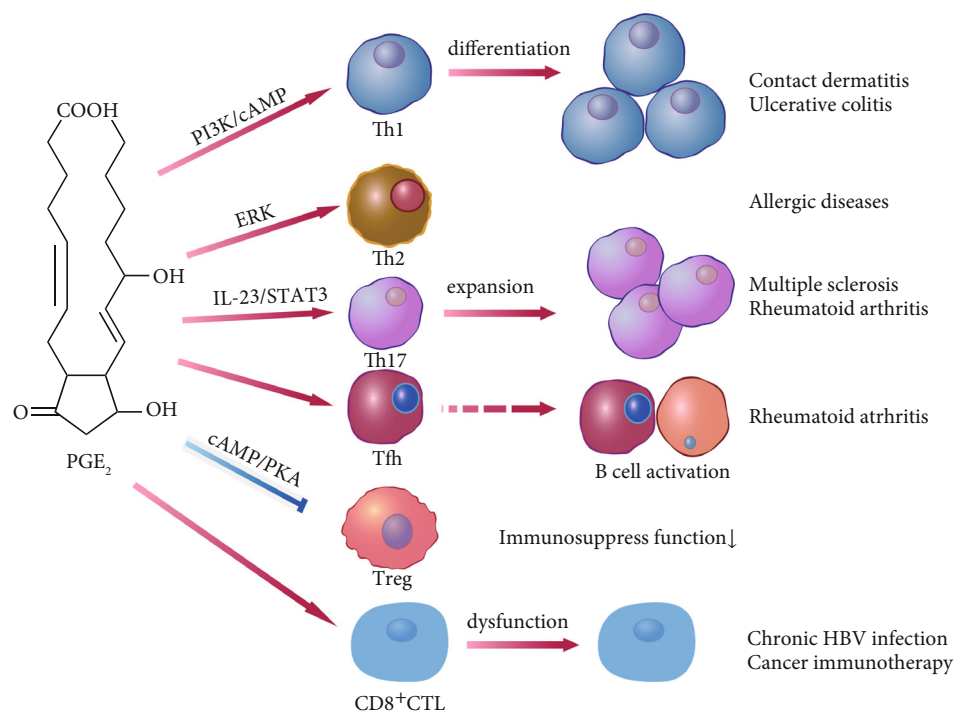


FIGURE 3: Regulation of T cell subsets by PGE₂. PGE₂ activates the differentiation of Th1 cells by decreasing the levels of cAMP; PGE₂ can enhance the IL-4/IFN- γ ratio in CD4⁺ T cell culture and regulates CD4⁺ T cell polarization towards Th2 cells; PGE₂ promotes the differentiation of naive CD4⁺ T cells into Th17 cells via the EP2 or EP4 receptor and cAMP pathway; PGE₂ may promote Tfh differentiation; PGE₂ suppresses the EP/cAMP/PKA pathway and inhibits Treg cell differentiation; PGE₂ inhibits the antiviral activity of CTLs and the survival of patients receiving cancer immunotherapy.

dysfunction correlates with PGE₂ levels in chronic hepatitis B patients. Patients with high levels of PGE₂ had more CD8⁺ T cells with remarkably low granzyme B expression and slightly low perforin expression; these proteins are two important effector molecules for the antiviral activity of CD8⁺ T cells in chronic hepatitis B patients [53]. Chen et al. showed that PGE₂ suppresses exhausted antigen-specific CTL function and promotes CTL apoptosis. The comodulation of PGE₂ and PD-1 signaling may represent a potent therapeutic avenue for the treatment of chronic viral infections [54]. Kim et al. demonstrated that the elevated mPGES1 expression is associated with low CD8⁺ T cell infiltration into melanomas and poor patient survival [55]. However, concrete results regarding the relationship between PGE₂ and CTL differentiation are still lacking. Notably, CD4⁺ T cells with cytotoxic activity (CD4⁺ CTLs) have been observed in various immune responses. These cells are characterized by their ability to secrete granzyme B and perforin and to kill target cells in an MHC class II-restricted fashion. CD4⁺ CTLs seem to be derived from various types of CD4⁺ T cells, and several differences have been observed during their differentiation [56]. Most likely, the differentiation of CD4⁺ CTLs can be influenced by PGE₂.

3. Conclusions and Future Perspectives

In this review, we highlight the immunoregulatory roles of the lipid mediator PGE₂ in controlling the differentiation of T cells. The dysregulation of T cell differentiation leads

to the occurrence of various diseases. Although PGE₂ exerts effects on different CD4⁺ T cell subsets, its role in Tfh cells and CD8⁺ T cells is not well known. The sophisticated interactions between PGE₂ and T cells still need to be further studied.

Various cells can secrete this lipid mediator, whereas all cells can generate responses to the signaling of PGE₂ through the EP receptors. Importantly, studying the signaling pathways of PGE₂ may contribute to further elucidating the pathogenesis of diseases, and targeting PGE₂ may be used to develop personalized therapies for RA, AS, neuropathic pain, tumors, inflammation, and so on. PGE₂ may be considered the point of intersection of different human systems. Thus, understanding the pathway of PGE₂ would yield novel insights for designing more effective therapies for immune diseases.

Conflicts of Interest

The authors declare that they have no conflicts of interest.

Authors' Contributions

XN conceived and designed the manuscript. YA wrote the manuscript. JY helped with manuscript preparation and literature search. YA, JY, and XN edited and critically evaluated the manuscript.

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