

REVIEWS: CURRENT TOPICS

Trapping of growth factors by catechins: a possible therapeutical target for prevention of proliferative diseases

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Abstract

The prevention of cancer through dietary intervention is currently receiving considerable attention. Several epidemiological studies substantiate that green tea has a protective effect against a variety of malignant proliferative disorders such as lung cancer, breast cancer and prostate cancer. This preventive potential of green tea against cancer is attributed to the biologically active flavonoids called catechins. Epigallocatechin 3-*o*-gallate, the major catechin found in green tea, mediates diverse physiological and pharmacological actions in bringing about the regression of the tumors and also lowers the risk of nonmalignant cardiovascular proliferative diseases. Much of the current research is being focused on how these catechins specifically bring about the regression of the experimentally induced tumors both in vitro and in vivo. These catechins exert diverse physiological effects against proliferative diseases by several mechanisms, most of which are not completely characterized. This review summarizes the mechanisms by which these catechins play an essential role in regulating the process of carcinogenesis, with a special emphasis on how these catechins antagonize the growth factor-induced proliferative disorders.

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1. Introduction

Tea, especially green tea, is the most consumed beverage in the world, next to water. Epidemiological studies demonstrated that the incidence of stomach cancer is low among tea drinkers in Shanghai [1]. It has also been reported that incidence of prostate cancer in China where green tea is consumed on a regular basis is the lowest in the world [2]. Similar studies reported that in Japan the onset of cancer in females who had consumed 10 cups of green tea per day was 8.7 years later and 3.0 years later among males, in comparison with patients who had consumed under three cups per day [3], and recurrence of stages I and II of breast cancer was reduced significantly by a daily consumption of four to five cups of green tea in Japan [4–6]. These findings prompted the world scientific community to unravel the startling anticancer properties attributed to green tea. Since then, much of the attention has been focused on finding a possible candidate against the

proliferative diseases through the green tea extracts. For the last two decades, extensive investigations have been carried out to figure out the green tea factor(s) that contribute(s) to the observed antiproliferative activity of the green tea. Until now, the results coming out with the green tea extracts are promising and raising hopes for finding out a possible antiproliferative agent for the cure and prevention of several proliferative disorders, especially the cancer. To transform the hope into reality that the factors from green tea are the possible effective antiproliferative agent, much of the research is being carried out worldwide and some of them are already in clinical trials [4–6]. The mechanisms by which these green tea extracts control the process of carcinogenesis and in the cancer prevention are diverse and these extracts employ several mechanisms, all at a time, to bring about the regression of the tumors in an efficient way. The antiproliferative effects of green tea catechins are partially explained by their antioxidative properties [7,8]. Recent investigations reveal that the green tea catechins exhibit several novel mechanisms to ward off proliferative diseases. Hence, this review focuses and portrays the novel and update mechanisms by which these green tea catechins prevent carcinogenesis,

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with a special focus on the newly proposed and proven concept that these catechins are the scavengers of the growth factors that promote carcinogenesis and other proliferative diseases.

2. Green tea

Green tea, a lipid extract obtained from the plant *Camellia sinensis*, possesses more pharmacological benefits when compared to black and oolong tea, even though they all are derived from the same plant but produced according to the different processes of drying and fermentation. “Black tea” is fully fermented, “oolong tea” is partially fermented and “green tea” is not at all fermented but only steamed. To prepare black and oolong tea, the young leaves are picked, allowed to wilt and then rolled. The leaves are then allowed to ferment, converting the polyphenols to phlobaphenes and forming aromatic compounds. During the fermentation process, the enzymes from leaf including polyphenol oxidase react with tannins and catechins. For the green tea preparation, the young leaves are not allowed to oxidize by fermentation but are steamed to inactivate the enzymes thereby preserving as much as 90% of the polyphenols contained in fresh leaves from being degraded [9]. The preventive potential of the green tea against cancer as evidenced from the several experimental studies is mainly due to these polyphenols, which are found in green tea in a relatively higher concentration compared to black and oolong tea, and hence the green tea possesses more anticarcinogenic potential in comparison to black and oolong tea. Of the total amount of tea produced and consumed in the world, 78% is black tea, 20% is green tea and less than 2% is oolong tea. Green tea is consumed mostly in China, Japan, India and a few countries in North Africa and the Middle East while black tea is consumed primarily in Western countries and in some Asian countries and the oolong tea in southeastern China and Taiwan [10].

3. Catechins, general cellular mechanisms of action

Dried tea leaves are composed mainly of phytochemicals known as polyphenols (30–36%), mainly flavanols (including catechins), flavonoids and flavandiols. The majority of the polyphenols are flavanols, more commonly known as catechins [11]. The main catechins found in green tea are catechin, epicatechin (EC), epigallocatechin (EGC), epicatechin-3 gallate (ECG), and epigallocatechin-3-gallate (EGCG) (Fig. 1). The catechin in green tea that has gained the most attention with respect to the anticarcinogenic activity is the potent antioxidant EGCG. Much of the anticarcinogenic effect of green tea is mainly attributed to EGCG. EGCG makes up about 10–50% of the total catechin content and appears to be the most powerful of all the catechins — with an antioxidant activity about 25–100 times more potent than vitamins C and E. EGCG has both

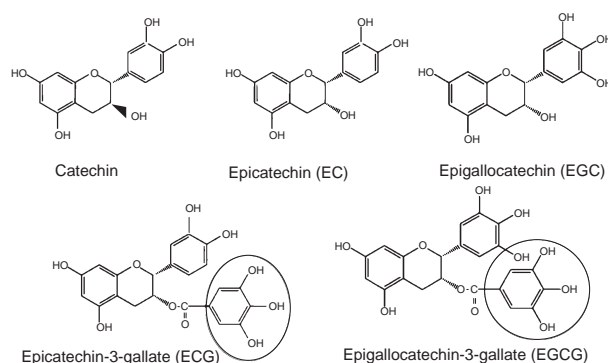


Fig. 1. Chemical structures of the green tea catechins. Only the catechins with the gallate group at 3' position are the ones having high efficacy in inhibiting the proliferation of cells in in vitro tumor models. The gallate moiety in these catechins is rounded off.

antimatrix metalloproteinase and antiangiogenesis activities [12,13]. In black tea, the major polyphenols are theaflavin and thearubigins.

The mechanism of action of catechins on warding off the cancer is diverse and involves several modes of actions. EGCG, the primary catechin in green tea, appears to inhibit the growth of cancerous cells as well as play a role in stimulating apoptosis, both of which are crucial aspects of cancer prevention [14,15]. EGCG inhibits the cellular proliferation primarily by

1. acting as antioxidants and scavenging the free radicals [16–18];
2. inhibiting the enzymes involved in cell replication and DNA synthesis [19–22];
3. interfering with cell-to-cell contact adhesion [12,23–26] and
4. inhibiting some of the intracellular communication pathways required for cell division [27–30].

The recent studies about the inhibition of cancerous growth/proliferation by catechins are revealing the novel modes of actions of catechins.

4. Effect of catechins on the growth factors-induced cell proliferation and signal transduction pathways

Growth factors are the extracellular signal proteins that promote growth, proliferation, differentiation or survival of cells in animal tissues. These signal proteins, also called mitogens, mostly act through receptor tyrosine kinases (RTKs, the second major type of cell-surface receptors next to G protein-linked receptors) located on the cellular membrane. The notable growth factors playing an essential role in stimulating the proliferation of various cell types are platelet-derived growth factors (PDGF AA, BB, AB), epidermal growth factor (EGF), fibroblast growth factors (FGFs: FGF-1 to FGF-24), hepatocyte growth factor, vascular endothelial growth factor (VEGF), insulin-like

growth factors (IGF-1 and IGF-2), macrophage colony-stimulating factor and all the neurotrophins, including nerve growth factor. The RTKs are transmembrane proteins with their ligand-binding domain on the outer surface of the plasma membrane. Their cytosolic domain either has an intrinsic enzyme activity or associates directly with an enzyme. To activate a RTK, the ligand usually has to bind simultaneously to two adjacent receptor chains. Platelet-derived growth factor, for example, is a dimer, which cross-links two receptors together. This cross-linking enables the neighboring kinase domains of the receptor chains to cross-phosphorylate each other on multiple tyrosines, a process referred to as autophosphorylation.

Autophosphorylation of the cytosolic tail of RTKs contributes to the activation process in two ways. First, phosphorylation of tyrosines within the kinase domain increases the kinase activity of the enzyme. Second, phosphorylation of tyrosine residues outside the kinase domain creates high-affinity docking sites for the binding of a number of intracellular signaling proteins in the target

cell. Each type of signaling protein binds to a different phosphorylated site on the activated receptor because it contains a specific phosphotyrosine-binding domain that recognizes surrounding features of the polypeptide chain in addition to the phosphotyrosine. Once bound to the activated kinase, the signaling protein may itself become phosphorylated on tyrosines and thereby activated. Alternatively, the binding alone may be sufficient to activate the docked signaling protein. Because different RTKs bind to different combinations of these signaling proteins, they bring about different cellular responses like cellular growth, proliferation and differentiation.

The growth factors usually act as local mediators at very low concentrations (about 10^{-9} to 10^{-11} M). The responses to them are typically slow (on the order of hours) and usually require many intracellular signaling events that eventually lead to changes in gene expression. These RTKs also have been found to mediate direct and rapid effects on the cytoskeleton, controlling the way a cell moves and changes its shape. These growth factors are not often

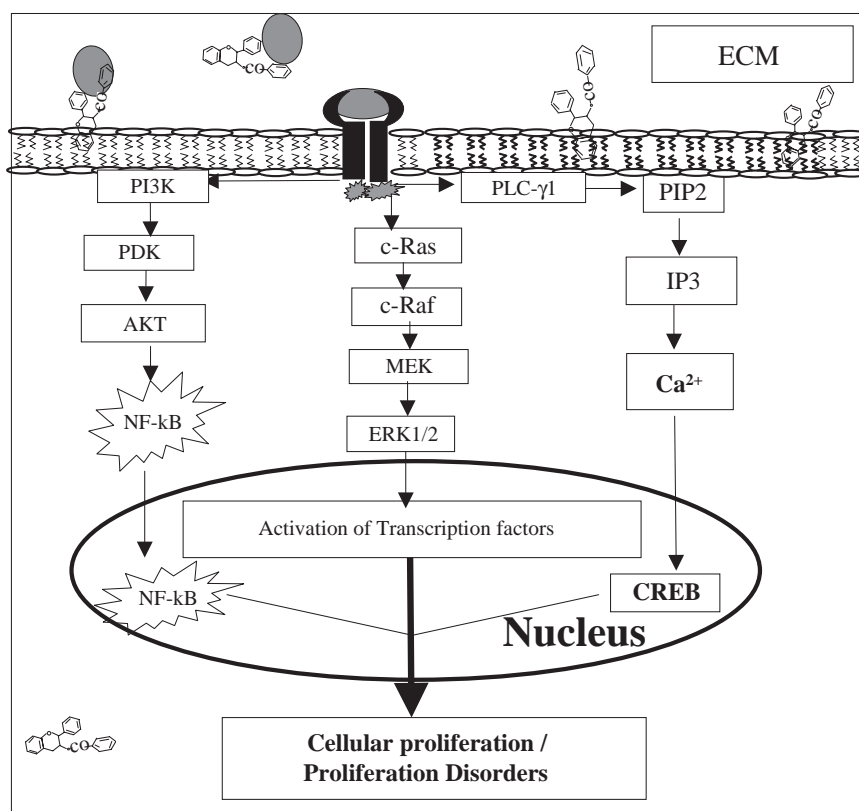
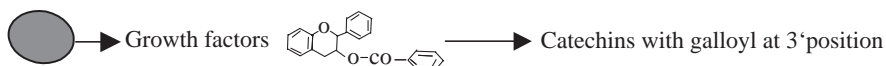


Fig. 2. Inhibition of growth factor-induced cell proliferation by EGCG and other catechins with a galloyl group at position 3. The growth factors transduce their mitogenic stimulus through RTK to activate the transcription factors such as ATF-2, Fos, Jun, Myc, and others. The catechins irreversibly bind with the growth factors. If the EGCG is already bound with membrane, then the EGCG-bound growth factors are immobilized on the membrane and thereby the growth factors are not available for binding with their respective cognate receptors. If the EGCG exists as free molecule, then EGCG-bound growth factor is free to move and may still mediate its mitogenic effect. Inhibition of intracellular signaling molecules by EGCG also is possible and reported.



diffusible but instead are found attached to surfaces over which the cell is crawling. Disorders of cell proliferation, differentiation, survival and migration are fundamental events that can give rise to cancer and abnormalities of signaling through RTKs have major roles in this class of disease. An increased activity of the RTKs such as the platelet-derived growth factor- α receptor (PDGF-R α), the β -receptor (PDGF-R β) and EGFR contribute to the development of proliferative diseases such as cancer and atherosclerosis [31–34].

Platelet-derived growth factor is a dimeric protein composed of A and B chains. The A and B chains are able to form three isoforms: PDGF-AA, PDGF-AB and PDGF-BB [35,36]. All three isoforms bind with high affinity to the PDGF-R α while only PDGF-BB binds with high affinity to PDGF-R β . PDGF-R α and PDGF-R β transmit the mitogenic signals through ligand-induced receptor autophosphorylation. Tyrosine-phosphorylated PDGF-R α or PDGF-R β induces tyrosine phosphorylation of different substrate proteins, including phospholipase C- γ 1 (PLC- γ 1), phosphatidylinosi-

tol 3'-kinase and extracellular response kinases 1/2 (ERK1/2) via the p21ras/mitogen-activated protein kinase pathway [37,38]. Activation of PLC- γ 1 results in hydrolysis of phosphatidylinositol 4,5-bisphosphate to diacylglycerol (DG) and IP3. The DG remains embedded in the plasma membrane and activates the Ca²⁺ protein kinase C. IP3 triggers the release of Ca²⁺ from endoplasmic reticulum and thereby increases the concentration of Ca²⁺ in the cytosol [39]. The increased levels of the cytosolic Ca²⁺ activate Ca²⁺/calmodulin-dependent protein kinases and eventually exert their effect on the transcription of the genes, thereby bringing about the cellular proliferation (mitogenesis). Autocrine activation of the PDGF-R β seems to be the initial cause of the development of the human A172 glioblastoma cells [40]. EGCG and other green tea catechins inhibit the proliferation of cells in several ways and their effects on the growth factor-induced signaling and subsequent cell proliferation are shown in (Figs. 2 and 3).

4.1. Effect of EGCG on the angiotensin II-induced signal transduction pathway

Key events in the development of atherosclerosis, hypertension- and angioplasty-induced restenosis are vascular smooth muscle cell (VSMC) hypertrophy and hyperplasia. Angiotensin II (Ang II) is a potent VSMC growth factor that is produced in higher levels during the pathogenesis of cardiovascular diseases [41]. Ang II-induced VSMC hypertrophy is defined as an increase in cell size and protein content without an increase in cell number and DNA replication. The signal transduction pathways underlying the Ang II-induced growth response involve the activation of mitogen-activated protein kinases, which activates AP-1. Emerging evidence suggests that endogenous reactive oxygen species (ROS) synthesis promotes VSMC hypertrophy and proliferation [42]. The inhibition of VSMC hypertrophy by EGCG is partly by inhibiting JNK signaling pathway by Ang II apart from the ROS scavenging activity of EGCG [43].

4.2. Effect of EGCG on the EGFR-mediated signal transduction pathway

EGFR's tyrosine kinase activation is believed to initiate multiple cellular responses associated with mitogenesis and cell proliferation. The overexpression of EGFR might produce a neoplastic phenotype. EGCG inhibits the autophosphorylation of EGFR by its ligand, EGF, and blocks the binding of EGF to its receptors [44,45]. EGCG also significantly inhibits DNA synthesis and protein kinase activities of EGFR [46].

4.3. Effects of catechins on the VEGF signal transduction pathway

There is a large body of evidence supporting a central role of angiogenesis in tumor growth and metastasis. Accordingly, the expression of VEGF, the most potent

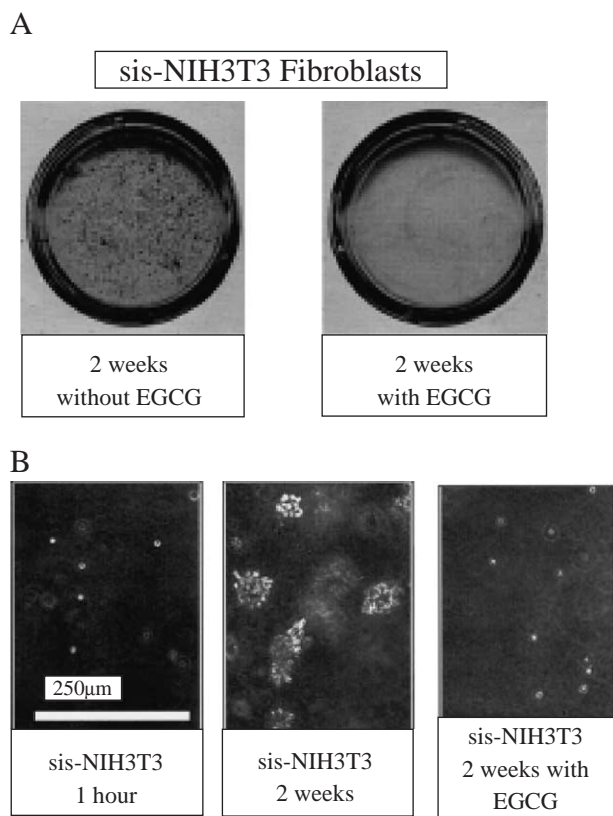


Fig. 3. (A–B) Anchorage-independent growth of *sis*-transfected NIH3T3 cells in the presence and absence of EGCG on semisolid agar. The oncogene *v-sis* of simian sarcoma virus is homologous to the cellular gene encoding the PDGF-B chain. Transfection of NIH-3T3 cells with *v-sis* leads to their neoplastic transformation because of the persistent autocrine activation of PDGF-B. The colony formation of *sis*-NIH3T3 cells was completely inhibited by EGCG. Reprinted from *Molecular Biology of the Cell* (Mol. Biol. Cell 1999 10:1093–1104) with the permission of The American Society for Cell Biology.

angiogenic stimulus known, and its receptors was found to be up-regulated in most of the tumors [47]. EGCG has been shown to inhibit growth and to induce apoptosis in different human cancer cell lines. It has been proposed that an anti-angiogenic property of EGCG may be one of the mechanisms leading to the inhibition of VEGF-induced carcinogenesis [45,48]. VEGF is a mitogen for endothelial cells that is often associated with tumor-induced angiogenesis [49]. VEGF binds to VEGF receptor-1 and -2, the latter being responsible for most of its mitogenic and chemotactic effects [49,50]. VEGF is involved in the angiogenesis of many solid tumors including breast cancer, colon cancer, hepatoma, bladder cancer, brain tumors and prostate cancer [51–57]. Moreover, there is increasing evidence that neovascularization is one characteristic feature of atherosclerotic plaques [58,59]. The two VEGF receptors form a dimer to activate autophosphorylation of tyrosine residues on the cytoplasmic domain [49]. The study by Neuhaus et al. [60] showed that treatment with EGCG results in an inhibition of human umbilical arterial endothelial cell (HUAEC) mitogenesis (DNA synthesis and cell proliferation). In addition, the signal transduction pathways of VEGF in HUAEC, including autophosphorylation of VEGF-R1 and -R2, phosphorylation of ERK1/2, as well as the expression of EGR-1 mRNA, are also inhibited in EGCG-pretreated cells [60]. Similar to “PDGF trapping by EGCG” as described below, trapping of VEGF by soluble and by membrane-anchored EGCG is likely and might play an important role in inhibiting the neovascularization process at the tumor site. In addition, it is well established that catechins are able to interact with several proliferation-related proteins [61–63]. Thus, the inhibition of growth factor binding to its receptor may represent a general principle in modulation of growth factor activity by EGCG. Taken together, the inhibition of VEGF-induced endothelial cell mitogenesis by EGCG, which persists after removal of the compound in addition to possible direct binding of growth factors, would result in a sustained inhibition of endothelial cell proliferation even at low plasma concentrations between doses. Thus, EGCG or other galloyl group-containing plant-derived catechins are attractive candidates for prevention and/or treatment of proliferative disorders.

4.4. The proposed EGCG receptor

Using a subtraction cloning strategy involving cDNA libraries constructed from cells treated or untreated with all *trans*-retinoic acid (ATRA) which enhances the binding of EGCG to the cell surface of cancer cells, the single target that allows EGCG to bind to the cell surface was discovered to be a 67-kDa laminin receptor (67LR) [64]. The 67LR is expressed on a variety of tumor cells, and the expression level of this protein strongly correlates with the risk of tumor invasion and metastasis. Characterizing the mechanisms by which EGCG acts through this 67LR should help in the design of new strategies to prevent cancer. The mechanism by which EGCG inhibits tumor growth via this 67LR has to be speculated in future, even

though it is not clear at present how EGCG mediates its antiproliferative effect via its receptor.

5. Trapping of PDGF by EGCG

Catechins with a galloyl group in the 3-position of the catechin structure, such as ECG, CG and EGCG, are able to suppress the PDGF-BB-induced stimulation of the PDGF-R β -mediated signal transduction pathways and mitogenesis of vascular smooth muscle cells [65]. The inhibitory effects of EGCG on PDGF-induced cell signaling and mitogenesis are due to the incorporation of EGCG into different cellular compartments, including cell surface membranes which leads to an irreversible and nondisplaceable binding of PDGF to nonreceptor binding sites, most likely resulting in a reduced PDGF binding to the respective receptors. Although only about 2% of the incorporated EGCG is found in cell surface membranes, the morphological analysis of EGCG-treated human VSMC revealed that most of the cytoplasmic EGCG is stored in vesicle-like inclusion bodies. The EGCG incorporated into these structures apart from the membranes is not likely to inhibit PDGF-R kinase or other signaling molecules [66]. This favors a surface membrane-linked mechanism of action of this catechin. In addition, the inhibitory actions of EGCG occur at a very early step in the PDGF signal transduction, namely, at the site of PDGF-R autophosphorylation. The incorporation of the EGCG onto the membrane has not altered the physical properties of the membrane as evidenced in terms of the membrane fluidity and hence the membrane embedded with the EGCG is intact without any distortion of the membrane structure [66]. Moreover, no inhibition by EGCG is encountered when soluble EGCG binds directly with the PDGF-BB. Hence, the inhibitory effect of EGCG on the PDGF-R β -mediated mitogenesis is mainly by “trapping” of the PDGF ligand by only the EGCG catechin incorporated or adsorbed onto the plasma membrane, which might have prevented the specific binding of PDGF-BB to its respective receptors. This means that the PDGF bound to free EGCG (i.e., non-membrane-bound EGCG) is still freely moving and in fact is able to cross-link and activate the specific receptors in the normal way as the free PDGF does. When the EGCG adsorbed on to the membrane binds irreversibly to the PDGF ligand, the resulting ligand–EGCG complex is immobilized on the membrane. Since the EGCG is anchored firmly on/in the membrane, the transducing potential of the bound ligand is abolished since the bound ligand cannot move freely to get in to contact with the specific receptors and cross-link them. That is, the ligands are “entrapped.” This mode of molecular action of EGCG provides an attractive model to explain the antiproliferative effects of the catechins [66]. It has been reported that all *trans*-retinoic acid (ATRA) enhances the binding of EGCG to the surface of the cancer cells [64].

An interesting aspect of the inhibition of PDGF-induced proliferation by EGCG is the preferential inhibition of PDGF-BB- as compared to PDGF-AA-induced cell signaling and mitogenesis. Some specificity for PDGF-BB is observed in human VSMC [66]. For example, EGCG only partially inhibits PDGF-AA-induced phosphorylation of PDGF-R α , as opposed to an almost complete inhibition of PDGF-BB-induced receptor phosphorylation. This apparent selectivity is supported by experiments with porcine aortic endothelial cells (AEC) stably transfected with PDGF-R α [66]. In these cells, EGCG did not inhibit PDGF-AA-induced phosphorylation of PDGF-R α , PLC- γ 1 or ERK-1/2. Similarly, PDGF-AA-induced $[Ca^{2+}]_i$ transients were not affected by EGCG. In contrast, EGCG markedly inhibited PDGF-BB-induced receptor phosphorylation and intracellular signaling events. In porcine AEC stably transfected with PDGF-R β , EGCG inhibited both PDGF-BB- and PDGF-AB-induced phosphorylation of PDGF-R β and PLC- γ 1. Thus, the specificity of the inhibitory effects of EGCG for the PDGF-BB-induced cell signaling is not absolute and not necessarily restricted to PDGF. Some degree of specificity for the PDGF-BB isoform is also observed for the inhibition of cell proliferation by EGCG. Although more pronounced effects are seen with PDGF-BB as a stimulus, cell proliferation induced by any PDGF isoform is inhibited by EGCG. Furthermore, in PDGF-R α -transfected porcine AEC, EGCG significantly inhibited PDGF-AA-induced mitogenesis without affecting PDGF-AA-induced signaling events (phosphorylation of PDGF-R α and of PLC- γ 1, $[Ca^{2+}]_i$ transients). Thus, the inhibitory effect of EGCG must be mediated by additional mechanisms independent of the early signal transduction pathway of the PDGF-R α , for example, by modulation of the expression/activity of cell cycle components [67].

Furthermore, a binding of PDGF to membrane components, such as gangliosides GM1 or GM2, has been reported [68]. These gangliosides have also been shown to bind to other growth factors, such as basic FGF (bFGF) [69]. Thus, trapping of growth factors by nonreceptor binding sites may represent a general principle in modulation of growth factor activity. Also, it is interesting to note that catechins not only bind to PDGF but are also able to interact with several other proliferation-related proteins [39,70–75].

6. Conclusion

Recently, extensive investigations have been made to develop synthetic RTK inhibitors as drug targets for therapy of cancer and cardiovascular diseases [76–78]. In this context, there are currently a large number of small-molecule RTK antagonists directed to PDGF-R, EGF-R and VEGF-R in phase I–III clinical development stages. In addition to direct binding of PDGF (and possibly other growth factors), the noncompetitive inhibition of PDGF-induced (also other growth factors like VEGF, EGF etc.)

mitogenesis by membrane-bound EGCG results in a sustained inhibition of cell proliferation even at low plasma concentrations. The mechanism of the inhibitory actions of EGCG, as proposed from our present studies, will help to better understand the beneficial effects of catechins in human epidemiological studies and in animal atherosclerosis models. Thus, EGCG or other galloyl group-containing plant-derived catechins are attractive candidates for the prevention of proliferative disorders.

Acknowledgments

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