Inflammation and cancer interconnection; simply as we think

Ahmed M. Hussein1*, Zeinab E. Darwish2 and Omar H. Soliman3

Abstract
Microbiota variety is site specific depending on its location in the body and this diversity can get together with human health. At the last decades, inflammation is a primary protective response that sometimes goes away and becomes a main cofactor in the pathogenesis of numerous chronic human diseases, including malignant tumor. As well, microbes have the potency power to influence tumor growth and progression through a wide forms of routes; including alteration of tumor microenvironment, prolonged activation of inflammation, induction of genotoxic restraints and metabolism. In this review, we will set a general overview of inflammation has suspected to provide a major role in the pathogenesis and development of cancer as an important object for advanced cancer biology.

Keywords: Inflammation, Cancer, Cytokine, Tumorigenesis

Introduction
The human body shortly becomes inhabited by microorganisms soon at birth. Microbes colonize zones that are immediately exposed to the air and surroundings including the nostrils, oral cavity, skin, stomach as well as the gastrointestinal and urogenital tracts. There is a large variation in the bacterial composition of sites within each organ between individuals. This variability is influenced by genetics, diet, medications intake and other external environmental factors. Additionally, the immune system affects species and localization of microbiota through congregate regulation of immune tolerance and inflammation [1]. The neoplasm has its microenvironment and inflammation; represent one of the hallmark of cancer and affect therapeutic resistance. The tumor immune response is inactiveequilibrium between antitumor mechanisms which serve to loweringthecancer growth and the pro-tumor inflammatory response which raising the immune tolerance, cell survival and reproduction. Inflammatory mediators exert pleiotropic actions in the growth of cancer. On one hand, inflammation advocacy carcinogenesis, malignant transformation, tumor growth, invasion and metastatic spread; on other hand, inflammation can activate immune effect or mechanisms that might limit tumor proliferation [2].

Based on the presence of leukocytes in cancerous lesions, Virchow made the initial connection between inflammation and cancer by monitoring leukocytes in neoplastic tissues [3]. During the last decade that is clear evidence of underlying molecular mechanism has been obtained suggesting that inflammation plays an important role in tumorigenesis and that chronic inflammation raising cancer risk. More than 25% of all of human cancer are related to prolonged inflammation and to viral and bacterial infections. However, the action of inflammation is not restricted to its role at tumor initiation and growth; inflammation can also be induced in growing tumor or as a response to anticancer therapy and cell death [4]. Table 1 provides an over view on inflammatory and pathogenic conditions that are considered to be associated with malignant transformation.

To know the role of inflammation in the development of cancer, it is important to understand what inflammation mean and how it give a share in physiological and pathological processes. Inflammation is portion of the complex biological response
of body tissues to harmful exciters such as pathogens, dead cells or irritants and is a covering response involving immune cells, molecular mediators and blood capillaries [3]. The role of inflammation is to remove the initial cause of tissue injury, eliminate necrotic cells from the original injury and to initiate tissue repair. The traditional signs of inflammation responses are swelling, heat, pain, redness and lack of function. Inflammation is a general response that is theorized as a mechanism of innate immunity, as compared to adaptive immunity; which is special for each pathogen. Minor inflammation could lead to advanced tissue damage by the harmful stimulus (e.g. bacteria) and disclose the survival of the organism. In contrast, prolonged inflammation may result in a host of diseases like chronic rheumatoid arthritis, atherosclerosis, chronic periodontitis and even malignancy (e.g. gallbladder carcinoma). Therefore, inflammation is normally carefully adjusted by the human body [5].

Inflammation can be classified as either acute or chronic. Acute inflammation is the early restraint of the body to foreign stimuli and is achieved by the increased escape of plasma and leukocytes (especially granulocytes) from the blood capillaries into the damaged tissues. A collection of biochemical events activates and matures the inflammatory response, involving the immune system, the innate vascular system and numerous cells within the damaged tissue. Prolonged inflammation; recognize as chronic inflammation leads to an advanced shift in the cells types present at the zone of inflammation and is characterized at the same timely destruction and curing of the tissue from the inflammatory responses. The inflammatory response must be actively end when no longer needed to stop the unnecessary damage to tissues. Failure to do that; results in prolonged inflammation and tissue damage [6].

Varies types of inflammation differing by cause, mechanism, result and strength can promote cancer growth and development. Chronic inflammation associated with infections or autoimmune disease foregoes tumor growth and can contribute to it during induction of oncogenic mutations, genomic instability, initial tumor promotion and enhanced angiogenesis. Chronic exposure to environmental irritants can lead to low grade chronic inflammation that precedes tumor development [7]. Several type of cancers developed from sites of infection, chronic inflammation and irritation. It is becoming clear that the tumor microenvironment which is largely coordinated by inflammatory cells is a fundamental sharer in the neoplastic process, proliferation, survival and migration. In addition, cancer cells have some of the signaling molecules of the innate immune system such as chemokines, selectins and their receptors for infiltration, migration and metastasis spread. These ideas are fostering new anti-inflammatory therapeutic approaches to cancer growth and development [8]. Tumor associated inflammation goes together with cancer development. This inflammatory response can foster new angiogenesis, rise tumor progression and metastatic expansion cause innate immunosuppression and further genomic instability. Cancer treatments can also move to cause an inflammatory response by causing trauma, necrosis and tissue damaged that stimulate tumor recurrence and resistance to therapy. However, in some cases, treatment induced inflammation can promote antigen presentation result in immune mediated tumor enucleation [9] (Figure 1).

**Table 1. Examples of Inflammatory Conditions that are Associated with Malignancy [4].**

<table>
<thead>
<tr>
<th>Inflammation that causes insults or pathological conditions</th>
<th>Associated malignancy</th>
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<tbody>
<tr>
<td>Silica, asbestos, smoking-associated siliosis and bronchitis</td>
<td>Lung carcinoma</td>
</tr>
<tr>
<td>Pelvic inflammatory disease</td>
<td>Ovarian carcinoma</td>
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<tr>
<td>Chronic indwelling urinary catheter</td>
<td>Bladder carcinoma</td>
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<tr>
<td>TRYPI mutation-associated pancreatitis and alcoholism-associated pancreatitis</td>
<td>Pancreatic carcinoma</td>
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<tr>
<td>UV irradiation-associated skin inflammation</td>
<td>Melanoma</td>
</tr>
<tr>
<td>Asbestos</td>
<td>Mesothelioma</td>
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<tr>
<td>Bile acids</td>
<td>Cholangiosarcoma and colorectal carcinoma</td>
</tr>
<tr>
<td>Gastric acid-associated Barrett’s metaplasia and reflux oesophagitis</td>
<td>Oesophageal carcinoma</td>
</tr>
<tr>
<td>Gall bladder stone-associated cholecystitis</td>
<td>Gall bladder carcinoma</td>
</tr>
<tr>
<td>Lichen sclerosis (a skin condition)</td>
<td>Vulvar carcinoma</td>
</tr>
<tr>
<td>Inflammatory bowel disease</td>
<td>Colorectal carcinoma</td>
</tr>
<tr>
<td>Hashimoto’s thyroiditis (an autoimmune disease of the thyroid) and Sjögren’s syndrome (an autoimmune disease of exocrine glands)</td>
<td>Mucosa-associated lymphoid tissue lymphoma</td>
</tr>
<tr>
<td>Gingivitis (inflammation of the gum tissue) and lichen planus</td>
<td>Oral squamous cell carcinoma</td>
</tr>
<tr>
<td>Sialadenitis (inflammation of the salivary gland)</td>
<td>Salivary gland carcinoma</td>
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**Review**

**Inflammatory Cell Component of Tumors**

Tumor cells produce numerous cytokines and chemokines that catch leukocytes. The inflammatory responses of a developing tumor may include a different leukocyte category. As a result
of these diversity types of inflammation, the cancer microenvironment has innate immune cells (including neutrophils, macrophages, dendritic cells, mast cells, myeloid-derived suppressor cells and natural killer cells) and adaptive immune cells (B and T lymphocytes) in addition to the cancer cells and their embracing stroma which formed from fibroblasts, endothelial cells, pericytes and mesenchymal cells [10]. Table 2 summarized the role of most inflammatory cells in antitumor and tumor-promoting cancer.

Pathways Connecting Inflammation and Cancer

The interaction between tumorigenesis and inflammation is mediated through intrinsic and extrinsic pathways. The intrinsic pathway is achieved by genetic alterations result in inflammation and neoplasia. These variations include mutation driven proto-oncogene activation, chromosomal rearrangement or amplification and inactivation of tumor suppressor genes. Transformed cells secrete inflammatory mediators and thus produce an inflammatory microenvironment. The extrinsic pathway is paid by inflammation or infections that rise the risk for the evolution of cancer in organs at risk such as the prostate, pancreas, colon, lung and skin [11]. The two pathways interfere in tumor cells and induce the activation of numerous transcription factors such as the nuclear factor (NF-κB), Signal transducer and activator of transcription 3 (STAT3), and Hypoxia-inducible factors (HIF-1) that lead to the formation of pro-inflammatory operators including chemokines, cytokines and Prostaglandin-end peroxide synthase 2 (PGHS-2). These molecules induct and activate various leukocyte populations such as macrophages, mast cells eosinophils and neutrophils into the tumor microenvironment like stromal and endothelial cells as well as infiltrating cells. This combined action of tumor and micro milieu results in a more pronounced generation of inflammatory mediators that drives the progression of appositive amplification loop which further triggers tumor infiltration and proliferation [12] (Figure 2).

Impact of Inflammation in Tumorigenesis

In the early stage of tumor development; inflammatory mediators such as cytokines, reactive oxygen species (ROS) and reactive nitrogen species (RNS) obtained from tumor

Table 2. Roles of Different Subtypes of Immune and Inflammatory Cells in Antitumor Immunity and Tumor-Promoting Inflammation [9].

<table>
<thead>
<tr>
<th>Cell Types</th>
<th>Antitumor</th>
<th>Tumor-Promoting</th>
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<tr>
<td>Macrophages, dendritic cells, myeloid-derived suppressor cells</td>
<td>Antigen presentation; production of cytokines (IL-12 and type I IFN)</td>
<td>Immunosuppression; production of cytokines, chemokines, proteases, growth factors, and angiogenic factors</td>
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<tr>
<td>Mast cells</td>
<td>Production of cytokines</td>
<td></td>
</tr>
<tr>
<td>B cells</td>
<td>Production of tumor-specific antibodies</td>
<td>Production of cytokines and antibodies; activation of mast cells; immunosuppression</td>
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<tr>
<td>CD8+ T cells</td>
<td>Direct lysis of cancer cells; production of cytotoxic cytokines</td>
<td>Production of cytokines?</td>
</tr>
<tr>
<td>CD4+ Th2 cells</td>
<td></td>
<td>Education of macrophages; production of cytokines; B cell activation</td>
</tr>
<tr>
<td>CD4+ Th1 cells</td>
<td>Help to cytotoxic T lymphocytes (CTLs) in tumor rejection; production of cytokines (IFNγ)</td>
<td>Production of cytokines</td>
</tr>
<tr>
<td>CD4+ Th17 cells</td>
<td>Activation of CTLs</td>
<td>Production of cytokines</td>
</tr>
<tr>
<td>CD4+ Treg cells</td>
<td>Suppression of inflammation (cytokines and other suppressive mechanisms)</td>
<td>Immunosuppression; production of cytokines</td>
</tr>
<tr>
<td>Natural killer cells</td>
<td>Direct cytotoxicity toward cancer cells; production of cytotoxic cytokines</td>
<td></td>
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<tr>
<td>Natural killer T cells</td>
<td>Direct cytotoxicity toward cancer cells; production of cytotoxic cytokines</td>
<td></td>
</tr>
<tr>
<td>Neutrophils</td>
<td>Direct cytotoxicity; regulation of CT1 responses</td>
<td>Production of cytokines, proteases, and ROS</td>
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infiltrating immune cells produce epigenetic variation in pre-malignant lesions and silence tumor suppressor genes. During tumor promotion, immune cells excrete cytokines and chemokines that work as survival and proliferation factors for cancer cells. The angiogenic transition is critical for an enough supply of tumor cells with nutrition, oxygen, growth and survival factors. During tumor progression and metastasis, both tumor and immune cells produce cytokines and chemokines result in an increase in cell survival, motility and invasiveness [13] (Figure 3).

**Inflammation and Tumor Initiation**

In the process of tumor initiation; normal cells acquire the initial mutational multiply that sends them on the tumorigenic track by providing development and survival priority over their neighbors. In most cases, however, a single mutation is scanty and many cancers need at least 4 or 5 mutations; it has been approached that an inflammatory microenvironment can rise the mutation rates and activating the propagation of mutated cells. Activated inflammatory cells act as origins of ROS and RNS that are able to producing DNA damage and genomic instability [15]. On the other hand, inflammatory cells may have cytokines like tumor necrosis factor (TNF-α) to stimulate ROS accumulation in adjoining epithelial cells. Inflammation produce mutagenesis may lead to inactivation or suppression of mismatch repair response genes, and ROS can also result in direct oxidative inactivation of mismatch repair enzymes [16]. Another mechanism linking inflammation with oncogenic mutations is up-regulation of activation produced cytidine deaminase (AID), an enzyme which promotes immunoglobulin gene class switching by catalyzing de-amination of DNA. The AID is overexpressed in many malignancies of diverse origins and its action is induced by inflammatory cytokines which produces genomic instability and rises mutation chance during error-prone ligation of double-stranded DNA breaks. Inflammation has been linked to epigenetic reprogramming which is encoded by an NF-kB target gene [17]. Other mechanism through which inflammation may activate tumor initiation is the induction of growth factors and cytokines that can allow a stem cell like phenotype upon tumor precursors or stimulate stem cell expansion, thereby connected to both stem cell reprogramming and stem cell renewal [18].

The interconnection between inflammation and cancer initiation is not a one-way street and there is also evidence that DNA damage can result in inflammation and thereby enhance tumorigenesis (Figure 4). One of the best examples is provided by the model of hepatocellular carcinoma induced by the carcinogen diethyl-nitrosamine, in which DNA damage contributes to necrotic cell death, resulting in an inflammatory action that activates tumor growth [19].
Inflammation and Tumor Promotion

Tumor promotion is the procedure of tumor development from a single initiated cell into a well-formed primary tumor. Early cancer development depends on increased cell generation and decrease cell death, both of which are produced by inflammation-driven actions. In verity, many of the inflammation activity on cancer are done at the level of tumor promotion and celebrated as tumor promoters’ inflammation. Tumor promotion induced by inflammation may occur early or late in cancer growth and can result in activation of premalignant conditions that were still for years. The strategies through which inflammation affects tumor promotion are variant and in addition to rise proliferation and allow survival can also involve the so known angiogenic switch, which promotes small dormant tumor to has the nutrition needful for the next developmental phase [20]. Mechanisms of inflammation driven tumor promotion are discussed as following:

A. Tumor-Promoting Cytokine Signaling

Producing of tumor promoting cytokines by immune/inflammatory cells that promote transcription factors such as NF-κB, STAT3 and activator protein (AP-1) in premalignant tissues to produce genes that activate cell proliferation and survival is a main tumor promoting mechanism. The NF-κB and STAT3 activate genes that control cell survival, proliferation and development as well as angiogenesis, invasiveness, motility and chemokine/cytokine production [21]. Oncogenic transcription factors competencetoinduced pattern recognition receptors by complex of bacteria and viruses. Nonetheless, the action of the cytokines that are activated in response to danger-associated (DAMP) or pathogen-associated molecular mechanisms during cancer growth is more firmly established in raising the tumor promotion [22].

The numerous cytokines (IL-1, IL-6, IL-23, TNF) and transcription factors (AP-1, STAT3, NF-κB) are sensible for both inflammation and tumor development, they rule the path of pro-tumorigenic signaling that may be aimed to limit both tumor associated inflammation and tumor development. Pharmacological intervention with cytokine signaling reduces tumorigenesis as well as tumor development and serve as a foundation for preventive and therapeutic approaches. Altogether, producing cytokine by immune and inflammatory cells is an important tumor promoting mechanism that allows malignant cells with a persistent production of growth and survival signals in a primary hostile microenvironment. In most cases, tumor promoting cytokines role in a paracrine mechanism, yet variant forms of cancer cells make their own cytokines, as IL-6 to have the same effect [23] (Figure 5).

B. Inflammation and Angiogenesis

Development of massive tumors needs an elevation of intratumoral blood feeding. This is triggered by tumor hypoxia, that induce angiogenesis and develop the ability of metastasis spread. Addition to hypoxia, cancer angiogenesis depends on recruitment of tumor-associated macrophages (TAMs) which active hypoxic signals and in role induce chemokines and proangiogenic agents. Induction of TAMs precursors are largely client on angiogenic mediators like; angiopoietin-2 and vascular endothelial growth factor (VEGF). Production of new lymphatic vessels is organized by VEGF-C and VEGF-D, whereas VEGF-A allows the recruitment of monocytes, which produce lymphoangiogenesis. Also, VEGF-A produced by myeloid cells prevents pericytes maturation and endothelial coverage of recently formed blood capillaries, and its surgical ablation activation tumorigenesis [24] (Figure 6).

As most developing tumors have some zones of hypoxia, there is not evident whether hypoxia is the main operator of tumor angiogenesis or whether hypoxic activate producing inflammatory signals that induce angiogenesis. Inactivation of NF-κB or STAT3 or TAM depletion unequivocally results in disrupted angiogenesis and reduce tumor development that confirm the critical action of inflammatory mediators in cancer angiogenesis [25].

C. Target Genes that Mediate Tumor Promotion

Several of the genes that activate the tumor promoting func-
tions of NF-κB, STAT3 and AP-1 have not been wholly realized, and most likely the pro-tumorigenic hints of these transcription factors are induced through different effectors. Some aims may be stripped by more than one transcription factor and may be more important in one cell type than in another [26]. Another group of target genes which allow tumorigenesis are chemokines and cytokines that labor in autocrine or paracrine pattern to ensure the continuous needs of inflammatory cells into the cancer microenvironment. The persistence of chronic inflammation is mainly achieved through positive feedback loops, which include inflammatory cells producing cytokines that allow chemokine synthesis in malignant and stromal cells leading to chronic recruitment of inflammatory cells into the tumor microenvironment. In many cases, the definitive chemokines are not produced by malignant cells but are produced in tumor associated fibroblasts upon connection with malignant cells [27].

Inflammation and Metastasis

From a clinical aspect, metastasis is the most critical view of tumorigenesis, because more than 90% of malignant mortality is result from metastasis. Different searches show that metastasis needs close interaction between tumor cells, inflammatory and immune cells and stromal elements. The spread of cancer can be mainly divided into 4 major steps:

The first step is act by epithelial-mesenchymal transition (EMT), in which cancer cells have fibroblastoid features which chrise their motility and induce them to penetrate epithelial linings/basal membranes and invade the blood vessels or lymphatics [28]. In the second step, malignant cells intravasate through blood vessels and lymphatics. Inflammation may induce this through formation of mediators that allow vascular permeability. This is followed by the third step, in which metastasis beginning cells survive and spread throughout the circulation. It has been estimated that nearly 0.01% of malignant cells that invade the circulation will finally survive and produce micro metastases. Finally, single metastatic progenitors interconnected with immune, inflammatory and stromal cells and beginning to proliferate. Many of these cells may formerly be acted to the pre-metastatic niche in restraint to tumor generated inflammatory signals before to the access of metastasis initiating cells [25 (Figure 7).

The transforming growth factor (TGF-β) signaling is aseriosregulator of the EMT and metastasis, and elevated TGF-β is mostlylyinked to poor prognosis. In spite of the defects in TGF-β signaling such tumors can as yet metastasize. These resistance effects of TGF-β at variestates of cancergrowth await mechanistic explanation. Malignant cell infiltrationneeds extreme proteolysis of the extracellular matrix at the invasive area. Inflammatory cells are effective provenance of proteases that degrade the extracellular matrix. Once metastatic cells reach the circulation they required to live in suspension and prevent detachment induced cell death. The survival of circulating malignant cells is influenced by inflammatory mediators produced by immune cells in restraint to cancer derived or pathogen derived stimuli. In addition to NF-κB and STAT3 activation, numerous of these cytokines can physically connected cancer cells to TAMs, helping them to invade together throughout the circulation [30].

On other hand, single metastatic malignant cells, which are no longer sitting within an immunosuppressive area, may be aimed again by immunosurveillance. Exactly, in some cases, infiltration of tumors by activated T cells reduces the spread of metastasis. The connection of circulating malignant cells with platelets or macrophages may prevent them from NK cell-mediated killing, thereby overcoming immunosurveillance [31]. Systemic inflammation promotes connection of circulating malignant cells to hepatic sinusoids, and this process is ruled by neutrophil-dependent upregulation of adhesion molecules. Several pro-inflammatory cytokines that are elevated in the circulation of cancer patient upregulate expression of adhesion molecules on the endothelium or in target organs and thereby rise the probability of metastatic cell attachment. Somematatureform tumors contain infiltrates of diverse leukocyte subsets. Leukocyte complexity modify depending on the tissue or organ position and grades of malignancy [32], suggesting that immune based therapies will able to reflect these small difference and be more specialty.

Cell Players and Mechanisms

Myeloid cells

During homeostatic state, leukocytes are encumbrance with maintaining tissue health. Innate immune cells having macrophages, granulocytes, mast cells, dendritic cells, innate lymphocytes and natural killer cells, which represent the initial mechanism of defense versus pathogens and foreign bodies. In tumors, these facts miss to solve and therefore result inprolonged inflammation of the damaged neoplastic tissue. Chronically activated leukocytes provide direct and indirect growth factors that induce proliferation of malignant and stromal cells [33]. In addition, numerous leukocyte subsets predominantly macrophages, granulocytes, monocytes and mast cells producevariant classes of proteolytic enzymes that adjust the structure and function of extracellular matrix. So that
their chronic presence allows a survival benefits to improve cancer cells by preserving proliferative signaling, decreasing cell death in restraint to matrix detachment, production and maintenance of angiogenesis, facilitating cancer cell escape and weaken antitumor cytotoxic cell mediated killing of cancer cells [34].

**T cells**
The CD4+ T helper cells are the organizer of inflammatory procedure in tumor tissue. Anumberous T helper (TH) subsets (TH1, 2, 9, 10, 17 and 22) specialized for activation particular types of inflammation, induction through their excretion of a restricted set of cytokines enabling responses immunityto tailored the special pathogen encountered. All of these distinct CD4+ T cell types can give a share in tumorigenesis by different paths depending on context. For example, regulatory T immunosuppressive subset of TH cells prevent cytotoxic action of CD8+ T cells through preventing tumor rejection [35]. Although in generichavoring tumor refuse TH1 cells might contribute to tumor escape through secretion of interferon (IFN)–γ, which triggers expression of apoptosis ligand that provides off signals to cytotoxic T cells. Furthermore, selective evolutionary pressure by IFN-γ may result in tumor editing and selection of resistant clones, thereby facilitating tumor growth [36].

The IL-17 induced by TH17 cells can act with IFN-γ to allow secretion of the chemokines CXCL9 and CXCL10 by cancer cells which engage cytotoxic T cells. Such connectionactions of IL-17 and IFN-γ could possibly be able to cancer treatment. The immune checkpoint molecules such as apoptotic protein PD-1 (a T cell receptor that mediates T cell inhibition) and its ligands, PD-L1 and PD-L2 forms a main receptor/ligand prevention pathway regulating T cell responses. Expression of PD-L1 on surfaces of tumor cells and tumor infiltrating myeloid cells allow an off signal to cytotoxic T cells and thus induce tumor cells to escape immune-surveillance. Under continuous antigen exposure (such as in chronic infections or in tumor microenvironments) both CD4+ and CD8+ T cells upregulate PD-1 expression, contributing to T cell exhaustion. Blocking this pathway, such as, during prolonged viral infection, reinvigorates virus-specific CD8+ T cell responses and results in enhanced T cell effector responses and viral clearance. However, classical chemotherapy paradoxically rises the number of macrophages expressing PD-L1, through prevention CD8+ T cells and elevation the risk of therapy failure [37].

**B cells**
As the onlymaker of immunoglobulins, B cells are important for humoral immunity and also effect other leukocyte variant. For example, B cell induced paracrine factors can be causative and/or potentiate disease by sustaining chronic inflammation through autoimmunity [38]. The action of B cells in cancer is under greatresearch. In the skin, squamous carcinogenesis is restricted in the loss of B cells. Two mechanisms implicated in B cell dependent skin carcinogenesis: (I) When autoantibody IgG is deposited via neoplastic parenchyma through leaky blood vessels ligation of immune complex/Fcγ receptors on mast cells and macrophages fosters pro-angiogenic and immunosuppressive gene expression programs; (II) B cell secretion of IL-10 and TNFα activates pro-tumorigenic myeloid cells that allow foster cancer progression [39].

**COX-2**
The arachidonic acid (AA) cascade has a vital role in mediating either the suppression or production of the inflammatory response. The COX-2 is the initial regulatory enzymes responsible for the translation of AA into the several lipid mediators involved in many biological functions. COX-2 has noted tumorigenic advantages and contributes to carcinogenesis by allowing insensitivity to antigrowth hints, evasion of apoptosis, sustained angiogenesis and tissue invasion/metastasis. Overexpression of COX-2 has been associated with carcinogenesis in animal models, and in several human cancers. The indirect action of the COX-2/PGE2 pathway in adjusting the tumor immune microenvironment has also been suggested via IL-17 promoting macrophage segregation [40].

**NF-κB**
The NF-κB transcription factors are preserved as coordinating regulators of immune and inflammatory actions that have a vital role in oncogenesis. All NF-κB family are necessary for dimerization and binding to DNA elements. These dimers bind to inhibitory protein IkB family of proteins (inhibitors of NF-κB) stopping their binding to DNA domains and localizing them to the cytoplasm in most quiescent cells [41]. Furthermore, physical, physiological and/or oxidative stress lead to activation of innate immunological processes resulting in inflammation which is associated with activation of the NF-κB signaling pathway. The NF-κB has a twin effect on inflammation. On one hand, the activation of NF-κB, as portion of the acute immune response, activates cytotoxic immune cells against malignant cells [42]. However, the activation of NF-κB also lead to up-regulation of antiapoptotic genes and the induced expression of other pro-inflammatory cytokines (e.g., TNF-α, IL-1, IL-6 and IL-8) and adhesion molecules which result in the recruitment of leukocytes to the zone of inflammation. NF-κB activation is also engaged in growth regulation and contributes to tumor progression by controlling vascularization of tumors through upregulation of VEGF and its receptors. The activation of NF-κB causes an increase in expression of the transcription factor Snail, that is essential in the TNF-α-induction of EMT which enables tumor progression and metastasis [22].

**TNF-α**
The TNF-α is a key pro-inflammatory cytokine, secreted by inflammatory cells which is involved in inflammation as-

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sociated carcinogenesis. It was named TNF-α due to it can produce tumor regression via the inducement of cell death. TNF-α is concerned in inflammation and immunity; as well as in a multitude of biological processes globalprogramed cell death, cell survival, angiogenesis and tumor cell invasion and migration [43]. The TNF-α tumor promoting action is dependent on ROS and RNS which can produce DNA damage and allow tumorigenesis. Direct evidence also points to the action of TNF-α in the metastatic cascade. Administration of TNF-α result in significant increase of the number of lung metastases. So, although TNF-α is a cytokine with recognized anticancer feature that has been applied as an anticancer agent for the therapy of some patients havingmassive solid tumors, its compact as a constituent within a multipronged approach targeted at a broad-spectrum of aims will need to be carefully assessed in light of these complicated results [44].

### iNOS

During ideal physiological conditions, nitric oxide synthases NOS is induced by the constitutive forms of NOS (cNOS/eNOS) and modulates vitalcellular processes such as vasodilatation, cell survival and development. However, in chronic inflammatory cases the iNOS-NO role is upregulated, and quickly yields; NO-derived species with strong nitrosative advantages especially when other reactive species are as will produced. Once formed, NO-derived species can quickly interact with all cellular structures including proteins, lipids and DNA. Therefore, the mostly carcinogenic role of NO-derived metabolites is related to their ability to potentiate genomic instability as stimulated by the RNS peroxynitrite [45]. In malignant tumor, NOS can be provided from both host and cancer cells; therefore, blocking tumor-iNOS hold potential implications for healthy cells. The method of therapeutic delivery needs a degree of specificity for malignant cells [46].

### AKT

Protein kinases are an important house of regulatory enzymes needed for the development, division and proliferation of cells and they have been closely studied as possible mediators of oncogenesis. In particular, the kinase signaling pathway known as phosphatidylinositol 3-kinase/protein-kinase-B/mammalian target of rapamycin represents one of the intracellular cascades of utmost importance when examining cellular proliferation, differentiation as well as cytoskeletal reorganization. The dysregulation of this pathway can direct the cell towards a carcinogenesis. As well, evidence indicated that AKT allows NF-κB activation [47].

### CXC Chemokines

Chemokines were originally had ability to adjust the directional migration of leukocytes to inflammatory sites. This observation has role for tumorigenesis as inflammatory cell infiltration is a common feature of numerous cancers and has varied functional consequences. Chemokines or chemotactic cytokines are a group of small (8–14 kDa) heparin-binding proteins that interact with cognate cell surface receptors and actserious roles in a varies of physiological processes such as growth, host immunity and cellular trafficking. These functionally related small secreted proteins constitute the largest cytokine group in humans [48].

In addition to their role in cell migration and inflammation, the chemokine/chemokine receptor system impacts development and progression of malignant diseases by regulating tumor initiation, growth, survival, migration, adhesion, invasion, angiogenesis and metastasis. Chemokines and their receptors regulate tumorigenesis directly by acting on tumor cells, and indirectly by regulating the composition of the inflammatory infiltrate. The diversity of the chemokine/chemokine receptor system is such that it can both contribute to, and inhibit, key events relevant to the tumorigenic process [49].

### Future Directions

The variance between tissues in which inflammation clearly drives cancer development (such as the gastrointestinal tract, liver and lung) and organs that can has severe chronic inflammation that has no increased risk of malignancy (such as the joints in rheumatoid arthritis) may be in relation to the presence or function of commensal microbial populations. As well, regions that are characterized by intense epithelial–microbial interactions (such as the intestine, lung and the liver, which is constantly exposed to gut-derived microbial products) may show an inflammation induced tumor tendency that is distinct from tissues that feature near sterility (such as joints and the brain). Hence, the commensal microbiota may represent the hidden connection between inflammatory preconditioning and the danger of tumor development. Not all chronic inflammatory diseases rise the risk of cancer; on the contrary many of them such as psoriasis have been associated with a decrease incidence of cancer [50]. Furthermore, the microorganisms can influence the inflammatory processes that are associated with cancer development may lead a new phase in the treatment of cancer. In the future, the targeted elimination of cancer-associated microorganisms might provide a new therapy option which, if effective, seems very attractive because of its minimal expected side effects and the possibility of its preventive application.

### Conclusion

The concept of inflammation enhance cancer has been firmly established during the last decade but many of the pathways that connect inflammatory processes to their considerable modulatory effects on various stages of tumor development remain elusive. The great link between tumor cells and their inflammatory microenvironment have been specified and we have learned a mainsearch about the cell types and mechanisms that are implicated in the interplay of inflammation, genetic instability, neoplasia and chemotherapy. Nonetheless, fundamental questions remain to be answered.
and this limits our ability to ded the growing knowledge of cancer associated inflammation for the development of recent therapeutic approaches.

Competing interests
The authors declare that they have no competing interests.

Authors' contributions

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