

## REVIEW ARTICLE

### Enhancement of cancer immunosurveillance by infections: a particular hygiene hypothesis for tumor growth?

#### Author

Mohamed F. Mandour<sup>a,b</sup>

Jean-Paul Coutelier<sup>a</sup>

#### Affiliations:

<sup>a</sup> Unit of Experimental Medicine, de Duve Institute, Université Catholique de Louvain, 1200 Brussels, Belgium.

<sup>b</sup> Department of Clinical Pathology, Faculty of Medicine, Suez Canal University, Ismailia, Egypt.

#### Correspondence

Jean-Paul Coutelier

E-mail: [jean-paul.coutelier@uclouvain.be](mailto:jean-paul.coutelier@uclouvain.be)

#### Abstract

Cancer results from a multi-step biological process consecutive to uncontrolled replication of transformed cells in which interactions with the surrounding environment and the host immune system play a major role. Anti-tumoral immune responses, mediated mostly by cytotoxic T cells, natural killer cells, and NK/T cells are in charge for killing the malignant cells and eradicating the tumor. At the early stages of cancer development they usually provide the appropriate immunosurveillance that eliminates most of the transformed cells.

The connection between cancers and infections, mostly by viruses, has attracted major attention. Roughly 12% of all human cancers are caused by oncoviruses via complex mechanisms involving host genetic variability and viral oncogenesis, while, in contrast, oncolytic viruses selectively infect and kill malignant cells. In addition to these direct effects of viruses on tumor cells, infections by viruses as well as by other agents, as key activators of the immune system, may enhance the efficacy of cancer immunosurveillance through bystander modulation.

This review provides an overview of the concept of immunosurveillance with highlighting its main cellular arms. We discuss the role of infections on cancer development and especially evidences of a positive effect of infections on the inhibition of some cancer development through enhancement of innate immune responses. This effect of infection might constitute a peculiar type of hygiene hypothesis, which could lead to distinct frequency of some cancers in populations with different exposure to infectious agents.

## 1. Introduction

Cancer is the product of a multi-step biological process in which a single transformed cell leads to a clonal cancerous growth through consecutive divisions. This complex process involves different stages starting from initiation, which encompasses cell damage resulting in a permanently distorted growth potential, followed by progression, representing many rounds of cell replication mediating the gradual transition towards an independent, cancerous growth. Crucial expansion of these cancerous cells to distant sites resulting in numerous tumour sites has been recognized as metastasis [1].

Malignant tumor development generally occurs through a substantial period of life. Such a lengthy period is clear when comparing for example the starting age for smoking and the usual age at which diagnosis of lung cancer frequently happen [2,3] and is supported by many human and animal studies in which a variety of premalignant lesions has been recognized.

Both genetic and environmental factors are involved in this progression towards clinical cancer. Chemical carcinogens cause DNA damage in the exposed cells [4], leading to uncontrolled cell proliferation, sometimes just by producing mild toxic damage in affected tissue [5].

The immune system is a central actor in the outcome of cancer development [6–8]. Normally, a competent immune system prevents the development of emerging tumors, a concept known as cancer immunosurveillance. Indeed, cancer immunosurveillance functions as an efficient extrinsic tumor suppressor mechanism against transformed cells. This protection is mediated via diverse arms of non-specific (innate) and specific (adaptive) immune mechanisms [8].

## 2. Carcinogenesis and environmental factors.

In 1954, Armitage and Doll developed the concept of cancer multistage model involving several basic assumptions:

1. Malignant tumors ascend from sequential alterations of one progenitor cell.
2. The process of developing a malignancy is equally likely for all cells in the same tissue.
3. Malignancy development process in one cell is totally apart from the process in any other cell.
4. Once malignancy has developed in a cell, spread to an obvious cancer is fast and involves numerous cells in the same tissue, and possibly would include metastasis to another tissue [9].

### 2.1 Infections

Infections with bacterial, viral and parasitic agents might have of central role in cancer rise. Schistosomiasis is a widespread helminthic infection in Asia and Egypt. The eggs of *Schistosoma japonicum* or *S. haematobium* are deposited in the colonic and bladder mucosa, respectively, causing inflammation and subsequent colon or bladder cancer [10]. A liver fluke, *Opisthorchis viverrini*, infects millions of people in Thailand and Malaysia. The flukes settle in bile ducts and increase the risk of cholangiocarcinoma [10]. *Helicobacter pylori*, a common bacterial infection of the stomach is a major cause of gastritis, ulcers and gastric cancer [10]. Aflatoxin, a mutagenic toxin found in moldy peanut and corn products, have been shown to interact with chronic hepatitis infection resulting in liver cancer development [11- 13]. Various major viral infections have contributed to development of malignancy. These include hepatitis B virus, (HBV) hepatitis C virus (HCV), Epstein Barr Virus (EBV), high-risk Human Papillomaviruses (HPVs), Human T lymphotropic Virus-1 (HTLV-1), HIV and Kaposi's sarcoma herpesvirus (KSHV).

Globally, oncoviruses count for about 12% of human cancers [14-16]. Even with their high incidence, significance to public health, and liability to prevention and directed therapies, understanding virus-induced cancers still has difficult challenges attributed to inadequate animal models of the disease, different nature of virally provoked cancers, and the complex nature of the virus-host cell interactions leading to cancer development [17]. However, various underlying mechanisms have been hypothesized for cellular changes leading to cancer in the course of viral infections. They include (i) Associated chronic inflammation that drives reactive oxygen species production and initiates mutations. This is evident in chronic HBV and HCV infections, where virus induced inflammatory reactions may finally lead to hepatocellular carcinoma. [18]; (ii) Recognition of viral genomes or replicative intermediates by the host leading to induction of DNA damage response needed by many oncoviruses for their replication; (iii) Signaling mimicry by viral-encoded proteins that destabilize host signaling mechanisms regulating cell growth and survival. As a result, host cells gain genetic instability, which increases their mutation rate, and speed up acquirement of oncogenic host chromosomal alterations [19]. As the host develops immune defenses against invading viral infections, viruses themselves have evolved to escape this protection. Human oncoviruses develop also strong immune evasion strategies to establish chronic infections, including anti-apoptotic and proliferative programs that in turn provoke malignant features in the infected cell [17, 20].

### **3. Anti-tumoral immunity and immunosurveillance**

Despite being considered in the past a simple witness of the battle between pro- and antioncogenic signals, the immune system is currently being known as a central actor in the outcome of cancer development [21-23].

Immunity has apparently two contradictory effects on tumors. Normally, a competent immune system prevents the development of emerging tumors, a concept known as cancer immunosurveillance [23]. This assumption hypothesizes that the immune system can control tumour growth by identifying different antigens on cancer cell precursors and destroying them prior to becoming clinically apparent. Although only few data show immunological abolition of premalignant lesions *in vivo*, great evidence supports the cancer immune surveillance hypothesis [24]. For instance, endogenously produced interferon- $\gamma$  (IFN- $\gamma$ ) was shown to be protective against the growth of spontaneous, transplanted or chemically induced tumors by mouse treatment with neutralizing anti-IFN- $\gamma$  monoclonal antibodies [25, 26]. Alternatively, the immunological pressure induced by cancer immunosurveillance induces the intrinsic nature of developing tumors through immunoediting mechanism [27]. The 'Immunoediting' concept was established by the observation that tumors transplanted from an immune-deficient animal to a syngeneic immune-competent animal are often rejected by the recipient's immune system, while tumors rising in immune-competent animals generally grow unhindered after transplantation [28, 29]. This process includes three stages: elimination, equilibrium, and escape [30, 31]. Elimination phase corresponds to the classical assumption of cancer immunosurveillance, where transformed and early stage malignant cells are removed by immune cells. Equilibrium is the phase of immune-mediated latency following imperfect tumor destruction. Failure to eradicate all transformed cells results in the development of tumors with reduced immunogenicity that can escape immune destruction and even, control subsequent inflammatory responses to their own advantage.

In humans, severe primary immunodeficiencies are coupled with higher incidence of various cancers as lymphomas, stomach, breast, bladder and cervical cancers [32-35]. Moreover, high incidence of tumours associated with oncogenic viruses (HHV8-related Kaposi sarcoma, EBV-related Hodgkin's and non-Hodgkin's lymphoma, HPV-associated cervical cancer and HBV/HCV-related hepatocarcinoma) has been found in HIV-infected immunodeficient patients [36]. Remarkably, a CD4 T-cell count in peripheral blood of HIV-infected individuals is inversely associated with increased cancer risk for these type of tumors [37]. Similarly, immunocompetent mice are able to reject cancer cells expressing surface ligands that can activate natural-killer (NK) cells or cytotoxic lymphocytes [38, 39], whereas RAG2<sup>-/-</sup> mice lacking both T and B cells are more susceptible to spontaneous and carcinogen-induced tumours [40].

Over the recent decades, the role of components of the immune system like perforin [39], interferon- $\gamma$  [40] and lymphocytes [41] have been proven to limit the outgrowth of transplanted, carcinogen-induced, and spontaneous tumors. However, despite the fact that immune responses can defend against malignancy, other immune mechanisms (i.e. chronic inflammation) can support the initiation or development of cancer [42]. Both inherent and extrinsic immune-regulating mechanisms can affect and control tumor development and progression. For example, following chronic viral infections exhausted immune cells will affect anti-cancer immune responses [43].

Most solid tumors are infiltrated by a wide array of immune cells including T cells (both CD4 helper and CD8 cytotoxic T lymphocytes) and NK cells [44]. Although these infiltrating immune cells usually display inefficient anti-tumoral activity, the quality and magnitude of this infiltrate has been established as a prognostic indicator of disease progression

[45]. To induce an effector and memory T cell response, specific tumor antigens are required. MAGE-1 was the primary gene known to code a human tumor antigen that is recognized by T cells [46]. Now, many tumor-associated antigens (TAAs) have been defined. TAAs can be classified in 3 main groups: 1) tissue differentiation; 2) cancer-testis; and 3) naturally occurring over-expressed antigens [47]. Tissue differentiation antigens are shared antigens between tumors and the normal tissue of origin; (eg. Gp100, Melan-A/Mart-1, Tyrosinase) in melanomas [48-53] as well as prostatic specific antigen (PSA) in prostatic carcinoma [54, 55]. Many cancer-testis antigens have been identified and tested in clinical trials, including the MAGE-A1 [56, 57] NY-ESO-141 and SSX-2 [58]. Over expressed TAAs such as tumor suppressor proteins (e.g. p53), antiapoptotic proteins livin and survivin, hTERT, Mucin 1 (MUC1) have increased expression in tumor cells when compared to normal tissues [59-65]. Tumor-associated carbohydrate antigens (TACAs) are glycans uniquely expressed or over-expressed by tumors, correlating also with various stages of cancer development [66-68]. The tumor – specific nature of these neo antigens is advantageous for eliciting specific T-cell responses with no risk of autoimmune reactions. Cancer vaccines based on defined specific tumor antigens should indeed elicit a very specific effector and memory cell response [69]. In view of the large number of possible tumor antigens for each cancer, the use of whole tumor cell vaccination approach has been considered the most favorable policy to embrace all potentially relevant antigens. However, several strategies have been implemented aiming to improve immune responses to peptide-based vaccines, via provoking the innate immune response [70, 71]. Dendritic cell (DC)-based vaccines in which tumor antigens are loaded on DCs in the form of peptides, tumor lysates [72], or apoptotic debris [73], represent one of the most

promising strategies to achieve effective anti-tumoral responses [74, 75].

### 3.1 Mechanisms of immunosurveillance

The early stage of anti-tumor immune response, involve the stimulation of tissue-located innate immune cells, such as macrophages and neutrophils that generate pro-inflammatory cytokines and chemokines. This pulls other innate cells, including natural killer (NK) cells, which can recognize and kill directly transformed malignant cells. Later on, dendritic cells (DCs) can process tumor antigens produced by dying tumor cells and deliver those in the lymph node to naïve T lymphocytes, leading to the activation of antigen-specific cytotoxic T lymphocytes (CTLs) and helper T cells. These cells additionally assist in tumor destruction [76]. Nevertheless, innate immune cells cannot recognize canonical neo-antigens that arise during tumorigenesis unlike T lymphocytes. They engage innate receptors to recognize ubiquitous intracellular self ligands, such as nucleic acids, that stimulate responses in certain cancer-associated contexts. In addition, other innate receptors recognize ligands that are displayed primarily by abnormal cells, so-called “induced self ligands”

#### 3.1.1 NK cells

NK cells are the most effector arm of innate immunosurveillance of cancer that has been studied. Initially they were characterized by their strong ability to directly kill tumor cells *in vitro* without former recognition. Different mechanisms have been postulated for the tumoricidal action of NK cells. *In vivo* and *in vitro* studies have shown that perforin play a major role in direct tumor cell lysis [77-83]. Alternatively, the engagement of death receptor-mediated pathways like TRAIL and FasL can induce tumor cell elimination [80, 84, 85]. This capacity of NK cells to kill tumor

cells may explain their protective effect against cancer development [86-88]. NK cell infiltrates in tumor biopsies have been associated with better prognoses in cancer patients [89, 90]. Recently, enhancing the tumoricidal activity of NK cells became an interesting focus for therapeutic purposes. NK cells also modulate activity of other immune cells, such as dendritic cells and T lymphocytes, through cytokine secretion or various receptor-ligand interactions [91-91]. NK-cell-derived IFN- $\gamma$  has shown to polarize macrophages towards a tumoricidal “M1” phenotype that provide defense against carcinogen-induced sarcomas [26]. Cytokines secreted by innate immune cells can encompass additional direct tumoricidal activity. Stimulated NK cells are major sources for various cytokines including IFN- $\gamma$ , TNF- $\alpha$ . IFN- $\gamma$  in particular is hypothesized to be have potent antitumor effects, such as inducing MHC I expression and sensitizing tumor cells to CD8+ T cell killing. NK-cell-derived IFN- $\gamma$  is related with better survival of patients in some cancers [94]. On the other hand, TNF- $\alpha$  via triggering caspase 8-mediated apoptosis can have direct cytolytic activity against malignant cells [95]. Together, IFN- $\gamma$  and TNF- $\alpha$  can drive tumor cells into senescence [96].

#### 3.1.2 NKT cells

In the context of tumor immunosurveillance, NKT cells can kill malignant cells through direct cytotoxicity or via activation of other immune cells. Type I NKT cells can directly lyse tumor cells through perforin-dependent mechanism [97], an effect that can be potentiated by granzyme B [98]. High expression levels of tumour CD1d, which restricts response of NKT cells has been associated with lower metastasis rates [101], whereas tumour cells expressing CD1d display high *in vitro* and *in vivo* susceptibility to direct NKT cell lysis [99–103].

$\alpha$ -GalCer the first recognized NKT cell ligand, is a strong activator of type I NKT cells. In mice with B16 melanoma, the application of its synthetic form (KRN7000) prolonged their survival [104, 29]. Upon  $\alpha$ -GalCer stimulation of NKT cells a profuse quantity of IFN- $\gamma$  is released which is crucial for tumor protection [105, 106], including anti-metastatic activity in lung and liver metastasis models [107].

Type I NKT cells can also mediate tumor immunosurveillance through initiation of Th1 cytokine cascades. A strong correlation has been shown between the Th1 cytokine profile (in terms of IFN- $\gamma$ :IL-4 ratio or IFN- $\gamma$  production) and the extent of protection from tumor growth in CT26 lung metastasis mice model [108]. In different tumor models, IFN- $\gamma$  release by NKT cells is the best correlate for tumor protection [109-113]. Surprisingly, NKT cells can augment tumor immunity by shifting the action of immunosuppressive cells, as shown in a model of influenza A virus infection, in which the absence of type I NKT cells resulted in the expansion of myeloid-derived suppressor cells (MDSCs), which suppressed CD8<sup>+</sup> T cell immune responses [114].

Tumor-induced inflammation is usually accompanied by production of serum amyloid A1 (SAA-1) which increases the interaction between type I NKT cells and neutrophils. Type I NKT cells not only diminish the unfavorable effect of neutrophils by suppressing production of IL-10, and enhancing IL-12, but also re-establish proliferation of antigen-specific CD8<sup>+</sup> T cells [115].

However, Type II NKT may have an immunosuppressive role on tumor immunology through MDSC activation and production of suppressive cytokines such as IL-13 and TGF- $\beta$  [102, 116-119].

### 3.1.3 Macrophages

Whereas much data focus on the crucial roles for T- and NK-cells in tumor immune surveillance, little evidence can show that macrophages can kill malignant cells through phagocytosis [120]. The established consensus is that macrophage activity is mostly protumorigenic through their ability to encourage angiogenesis and metastases [121]. Tumor associated macrophages (TAMs) of the M2 type can, however, be re-educated back towards a tumoricidal M1 phenotype through the production of IFN- $\gamma$ , and the overexpression of miR-155 [27, 122].

### 3.1.4 Tumor-associated neutrophils

The presence of tumor-associated neutrophils (TANs) in human tumors correlates with advanced disease and poor outcome in several types of human cancer [123]. Although neutrophils usually play a role in tumor progression, their N2 phenotype can be reversed to an anti-tumoral N1 phenotype by TGF- $\beta$  blockade [124] or by the effect of IFN- $\beta$  [125].

### 3.1.5 Other innate cells of interest

Based on their cytokine profiles, transcriptional activity, and effector functions, innate lymphoid cells (ILCs) strongly resemble the different helper T cell subsets. Depending on the secreted cytokines and the specific tumor microenvironment, ILCs may either aid anti-tumor immune responses or promote tumor formation and growth [27, 126]. Gamma/delta ( $\gamma\delta$ ) T-cells may limit cancer incidence in skin cancer mouse models and in a transgenic model of prostate adenocarcinoma through direct lysis of tumor cells [127, 128].

#### 4. The hygiene hypothesis

The hygiene hypothesis, first proposed by Strachan in 1989 [129], suggests that the strong increase in the frequency of allergic diseases observed in advanced industrialized countries is caused by a reduced exposure to common infections in early childhood. Initially supported by epidemiological studies, a protective effect of infections on allergic diseases has been also reported in experimental models [130, 131]. It was first postulated that a decrease of Th1-inducing bacterial and viral infections would result in a Th2 immune microenvironment prone to the development of allergies. However, Th1 autoimmune diseases increase as well as allergies, and parasite-induced Th2 microenvironments do not trigger more allergies in their host [132, 133]. Rather than a shift in Th1/Th2 balance, it has then been proposed that repeated infections inducing pro-inflammatory responses would in turn trigger Treg cell activation and secretion of immunosuppressive cytokines, like IL-10 as a counter-regulation. Therefore, the increase in both allergic and autoimmune diseases would result from a reduced immune suppression that was previously a consequence of childhood repeated infections [132-134]. A role of gut microbiota composition has also been suggested as a factor triggering such a modulation of immune response and of resulting diseases [135]. Whatever the mechanisms involved, it seems established that bacteria, viruses and parasites may sufficiently modulate the host immune microenvironment to deeply change not only the course of concomitant diseases initially unrelated to the infection, but also the probability to develop immune-regulated diseases.

#### 4.1 Infection and cancer: a peculiar hygiene hypothesis ?

Could a decrease of infections similarly result in an increase in the development of some cancers ? As mentioned above, the relationship between infections and cancer has so far mostly been focused on the ability of several pathogens to trigger oncogenesis through cell transformation. This infectious agent-induced increase in the rate of cancers may sometimes result from inflammation, which would be a result radically different from what can be expected from an hygiene hypothesis.

Oncolytic viruses (OVs) can selectively infect malignant cells and kill them while sparing normal healthy cells. This selective oncolysis can be either (i) a natural feature of the virus, such as parvoviruses, myxoma virus or reovirus with minimal or no pathogenicity in humans; (ii) a product of genetically-engineered virus, with mutations/deletions in genes required for replication in normal, but not cancer cells. Those OVs include adenovirus, herpes simplex virus, and vesicular stomatitis virus [136, 137]. Moreover, OVs may induce an intense host immune response, leading to the damage of remaining malignant cells and lasting antitumor immunity. Several OVs provoke immunogenic tumor cell death (ICD, such as immunogenic apoptosis, necrosis, and pyroptosis, which activates host immune responses [138, 139]. ICD is associated by cell surface exposure of calreticulin and heat shock proteins and the production of some molecules like ATP, uric acid, and high-mobility group box 1 that possess immune-stimulating characteristics. Furthermore, ICD of tumor cells also liberates tumor-associated antigens that are crucial for generating an antigen-specific antitumor immunity [140-142].

Interestingly, a few clinical studies have reported an inverse relationship between an history of febrile infections or vaccinations and the development of melanoma [143-145], a

tumor that is known to be sensitive to destruction by NK cells [146]. In the mouse, acute infection with lactate dehydrogenase-elevating virus (LDV), a nidovirus that induces a strong modulation of the immune microenvironment of its host, including NK cell activation [147], prevents plasmacytoma growth [148]. A similar inhibition of cancer development after LDV infection has been observed with mesothelioma (Mandour, unpublished data). The protective effect of LDV infection depends on NK cell activation and on IFN- $\gamma$  production by those activated cells [148]. It may thus be postulated that repeated stimulation of the innate immune system, and especially of NK and/or NKT cells by infectious agents induce levels of IFN-  $\gamma$

sufficient to prevent the development of cancer cells sensitive to this cytokine. Such a protective effect of infections would constitute a peculiar type of hygiene hypothesis, that would be important to confirm in order to appropriately target preventive cancer diagnosis in populations with various levels of common infections.

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