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How Comorbidities Shape Cancer Biology and Survival

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Abstract

Comorbid chronic diseases affect cancer patients with an increasing frequency as populations get older. They negatively and disproportionately impact underserved populations and influence cancer diagnosis, tumor biology and metastasis, and choice of treatment. Many comorbidities are associated with a delayed cancer diagnosis. Although the relationship between comorbidities and cancer risk and survivorship has been studied extensively, we still lack knowledge on how they affect tumor biology and the metastatic process. Here, we will discuss our current understanding of mechanisms linking comorbidities to an adverse tumor biology and lethality and introduce thoughts of how we can close existing gaps in this knowledge. We argue that research into comorbidity-induced alterations in cancer metastasis, immunity, and metabolism should be prioritized.

Keywords

comorbidity; cancer; metabolism; immunity; metastasis; survival; obesity; diabetes

Chronic diseases modify cancer risk and survival

A comorbidity among cancer patients is generally defined as the coexistence of a disorder/ chronic disease in addition to cancer. These disorders include chronic cardiovascular, liver and renal diseases, diabetes, metabolic syndrome, connective tissue diseases, chronic infectious diseases, **dysbiosis** (see glossary), neurological disorders such as dementia and chronic stress and depressive disorders, and autoimmune diseases such as rheumatoid arthritis, inflammatory bowel disease, systemic lupus erythematosus, or Sjogren's disease. Although not a chronic disease, COVID-19 infections have recently been associated with an excessive mortality among cancer patients [1]. Many of these comorbidities share risk factors with cancer, thus commonly co-occur with cancer. It has been estimated that three-quarters of cancer patients have at least one comorbidity [2] while data from Medicare beneficiaries in the United States indicate a prevalence of 40% [3]. Comorbidities do not

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affect all segments of the US populations equally. Native Americans and African Americans have significantly elevated rates for obesity, diabetes, chronic kidney disease, and hypertension, when compared to other population groups [4].

Comorbidities can influence cancer outcomes. For example, epidemiology has shown that comorbid human immune deficiency virus (HIV) infections may increase cancer mortality [5]. They impede the participation of cancer patients in clinical trials and adversely affect the discussion and offer of trial participation [6]. The uptake of cancer screening is inversely associated with the severity of comorbidities, potentially delaying a cancer diagnosis in those most impacted by multiple comorbidities who also tend to be among the poor and underserved [2, 7]. In contrast, having just one comorbid condition may lead to increased screening participation because of increased contact with health services [8]. Yet, individual chronic diseases may have different effects on screening. Diabetes was found to associate with decreased cancer screening whereas an infection with HIV may not uniformly affect cancer screening decisions [9, 10]. Comorbidities have an adverse effect on cancer survival [3]. They are strong prognostic factors of poor survival in colorectal cancer patients independent of sociodemographic factors and tumor characteristics [11]. The negative impact of comorbidities on cancer outcomes tends to increase with increasing severity of the comorbidities and their impact is generally greater for cancers with otherwise better outcomes. The presence of a comorbidity will influence treatment selection and particularly the use of chemotherapy [12]. Cancer patients with a comorbidity are generally less likely to receive curative treatment for their cancer than those without the comorbidity [3].

In this opinion piece, we will summarize key evidence that links comorbidities to cancer development and an adverse tumor biology. We will then discuss the mechanisms that underly these relationships - to the extent they are known - and offer opportunities to close existing gaps in this knowledge. Much of the presented research investigated two comorbidities, obesity and diabetes, as modifiers of tumor biology and cancer outcomes because these conditions can readily be studied in experimental models of cancer. Other comorbidities still lack suitable experimental models or are studied with cancer models that may not fully recapitulate the human disease, such as the autoimmune component of inflammatory bowel disease.

Comorbid conditions and their treatment frequently impact tumor biology

Comorbidities may modulate cancer risks by affecting tumor biology (Figure 1). The hypothesis is robustly supported by clinical and epidemiological studies and laboratory investigations with animal models [2, 13]. Diabetes has been linked to a doubling of liver and pancreas cancer incidence and is associated with the risk of breast, cholangiocarcinoma, colorectal, endometrial and gallbladder cancer but may have a protective effect against prostate cancer [14, 15]. It is thought that diabetes and hyperinsulinemia promote cancer development and progression through insulin and insulin-like growth factor signaling, as indicated by human studies and further supported by experimental research [15–17]. Additional mechanisms, like chronic inflammation, are likely involved. Antidiabetic medications can interfere with tumor biology. Metformin inhibits breast, colorectal, and endometrial cancer development whereas other agents such as insulin (glargine),

pioglitazone, and sulfonylureas have been associated with modestly increased cancer risks [18]. A systematic review and meta-analysis established diabetes as a significant risk factor for non-alcoholic fatty liver disease (NAFLD) with an estimated 50–60% global prevalence of NAFLD among patients with type 2 diabetes [19]. Many of these patients will develop hepatic fibrosis, conferring increased liver cancer risk [20]. Patients with rheumatoid arthritis have a heightened risk of developing Hodgkin and non-Hodgkin lymphoma and the clinical course of these lymphomas is often aggressive [21]. The correlation between rheumatoid arthritis disease activity and lymphoma risk suggest that both chronic inflammation and immune stimulation induced by a rheumatoid arthritis are possible drivers for this relationship although experimental data supporting the hypothesis are missing. Alternatively, immune suppression induced by anti-rheumatoid arthritis therapy may increase the risk of developing lymphoma. Lastly, hypertension has been associated with an elevated risk of developing kidney, colorectal, and breast cancer [22]. Changes in calcium metabolism in hypertensive patients may broadly increase cancer cell proliferation while hypertension-induced chronic renal hypoxia, together with a deregulated renin-angiotensin system, is a proposed mechanism that may increase the risk for kidney cancer.

Comorbidities promote cancer metastasis

Comorbidities are associated with an increased risk of cancer-specific mortality, indicating that comorbid conditions may enhance the metastatic spread of cancer, the primary cause of cancer deaths [2]. Yet little is known about the mechanisms by which comorbidities may impact metastasis. Obesity frequently associates with lethal cancer [15]. In mouse models of breast cancer, a high-fat diet and obesity promoted metastasis through impaired tumor vascularization and increased mesenchymal differentiation of cancer cells, altered chemokine signaling, changes to the tumor immune environment, and activation of the sphingolipid pathway [23–25]. A mesenchymal differentiation of cancer cells is also induced by diabetes, as shown for non-small cell lung cancer patients [26]. **Hyperglycemia**, a hallmark of diabetes, increases metastatic seeding of 4T1 breast cancer cells. In the 4T1 mouse model of breast cancer metastasis, hyperglycemia impaired tumor vascularization and secretion of granulocyte colony-stimulating factor and subsequent recruitment of neutrophils into metastatic sites [27]. Others reported that hyperglycemia promotes metastatic colonization of pancreatic ductal adenoma cells through activation of the pro-metastatic Runx3/Col6a1 pathway [28], whereas obesity-induced inflammation may lead to pancreatic cancer progression and resistance to therapy through stellate cell activation and increased desmoplasia [29]. Heart disease and cancer have shared risk factors. Still, there is evidence that heart failure *per se* is oncogenic and promotes cancer development and spread through release of soluble factors like serpins and signaling molecules from the renin-angiotensin-aldosterone pathway [30, 31]. Psychosocial factors such as chronic stress and depressive disorders are associated with cancer survival [32]. A pro-metastatic niche has been described for breast tumors from socially isolated women [33] and a decrease in chronic depression may slow metastasis in breast cancer patients [34]. Chronic stress and sympathetic nervous system signaling enhance breast cancer metastasis in animal models through increased secretion of colony stimulating factor 1, recruitment of M2 macrophages, and vascular endothelial factor C-induced lymphatic remodeling, enhancing the odds of metastasis [35,

36]. Thus, there is indication that co-morbidities are causatively linked to cancer metastasis although the existing literature remains sparse and needs to be expanded. There is, however, a lack of studies exploring the biology of primary tumors that give rise to metastasis in cancer patients with co-morbidities, or of studies that investigate the characteristics of metastatic lesions in these patients, and how those may relate to co-existing comorbidities.

Comorbidity-induced changes to the tumor immune environment that drive cancer progression and therapy resistance

Various malignancies are reproducibly cured in mouse models using immune therapy and chemotherapy. Yet, most of these therapies show objective responses in only a fraction of treated patients in the clinic. One reason for this disconnect might be the use of young, lean mice that lack immune-altering comorbidities present in elderly cancer patients. Obesity, diabetes, and viral infections like HIV/AIDS or chronic viral hepatitis have been shown to cause significant changes to the immune system and may reduce local and systemic immunity. One study showed that genetically and diet-induced obesity in the B16 melanoma and 4T1 breast cancer models results in PD-1-mediated T cell dysfunction, increased immune aging, and tumor progression which was partly driven by leptin [37]. In this study, obesity was also associated with increased efficacy of **PD-1 blockade** in both tumor-bearing mice and cancer patients. Another study examined the effect of obesity in two immunotherapeutic models, namely systemic anti-CTLA-4 monoclonal antibody therapy and delivery of a TRAIL-encoding adenovirus plus CpG. Here, both therapies were effective in lean mice but did not provide a survival benefit in mice with diet-induced obesity. Further analysis showed that leptin was a mediator of therapy resistance in this model, suggesting that leptin is a therapeutic target to improve tumor immunotherapy when immune-modulating comorbidities are present [38]. Chronic stress and depression are other comorbidities with good evidence that they can affect the systemic and intratumor immune environment, namely through activation of hypothalamic-pituitary-adrenal signaling as the central stress response system [39, 40].

Obesity and diabetes are cancer risk factors that induce changes to the **gut microbiome** and increase the risk of mucosal infections, establishing a link between dysbiosis and cancer (Box 1). There is good evidence that obesity, diabetes, certain chronic infections, and stress exposures can alter the tumor immune environment and gut microflora, but most other cancer-associated comorbidities have not been studied in this context, and we still lack patient data that would confirm the findings from animal studies.

Chronic disease-induced inflammation is a cancer risk factor and modifier of tumor biology

Chronic infections commonly cause chronic inflammation which is a cancer risk factor. The link between chronic inflammatory diseases and cancer has been reviewed in the past [13]. Obesity causes macrophage accumulation in adipose tissue and systemic chronic inflammation, leading to obesity-related insulin resistance [53]. Both dietary and genetic obesity promote liver inflammation and tumorigenesis through upregulation of

proinflammatory cytokines, including interleukin 6 and tumor necrosis factor α [54]. Recently, it was shown that interleukin 6-induced androgen receptor signaling leads to up-regulation of cell cycle-related kinase and mTORC1-dependent metabolic and immunosuppressive reprogramming in obesity-associated hepatocellular cancer [55]. In breast cancer, obesity upregulates adipose inflammation and estrogen synthesis by the aromatase pathway and activates the NLRC4 inflammasome pathway leading to increased interleukin 1 β and VEGFA expression and enhanced tumor angiogenesis [56]. In pancreatic cancer, obesity-induced inflammation may lead to disease progression and resistance to therapy through stellate cell activation and increased desmoplasia [29]. Yet, many aspects of the inflammation-to-cancer axis remain incompletely understood for most obesity-associated cancer types. Furthermore, the role of inflammation in comorbidity-induced cancer progression has rarely been studied for comorbidities other than obesity and infectious diseases.

Impact of comorbidities on cancer metabolism

The metabolic health status of a person, rather than obesity *per se*, may confer cancer risk [57]. Comorbidity-induced changes in cancer metabolism have been studied for few conditions, namely, obesity, diabetes, metabolic syndrome, dysbiosis, and chronic stress exposure. Persistent metabolic alterations are hallmarks of cancer that maintain tumor growth and induce **epithelial-to-mesenchymal transition** and metastatic spread, but also provide vulnerabilities that make cancer metabolism a target for cancer therapy [58]. Stress-induced catecholamine signaling may promote cancer stem cell traits through a lactate dehydrogenase A-dependent mechanism that rewires cancer metabolism [59]. Reprogramming of cancer metabolism by aberrant Myc signaling or the acquisition of mutations in metabolic enzymes, such as isocitrate and succinate dehydrogenases, or fumarate hydratase, commonly affect mitochondrial metabolisms and may lead to excessive production of **reactive oxygen species**, further enhancing the mutational burden and malignant adaptations of cancer cells [58]. High-fat diet fuels prostate cancer progression by enhancing the Myc transcriptional program through metabolic alterations and histone methylation [60]. Hyperglycemia and diabetes increase protein kinase C signaling and formation of advanced glycation end-products and activate NF κ B. Activation of these and the hexosamine pathway is triggered by hyperglycemia-induced mitochondrial superoxide production and can be blocked through normalization of mitochondrial superoxide production [61, 62]. High-fat diet and obesity more generally alter lipid and cholesterol metabolism in adipocytes and cancer cells [63] and increase 27-hydroxycholesterol [64]. This metabolite is an endogenous estrogen receptor agonist and enhances metastasis in orthotopic breast and pancreatic cancer models, in part through its action on immune cells [64]. There is also evidence that obesity-associated pancreatic cancer is driven by metabolic alterations. A critical role of mitochondrial arginase, ARG2, and the urea cycle was identified using human pancreatic cancer cells and an orthotopic xenograft model of obesity-induced pancreatic ductal adenocarcinoma [65]. The data show a dependency on ARG2 for obesity-driven pancreatic cancer in this model. Lastly, colorectal and liver cancer patients are impacted by comorbidity-induced changes to the metabolism of the gut microbiome, as already discussed by us [45–47], highlighting the intertwined relationships between gut

microflora metabolism and cancer. However, findings from these primarily animal-based investigations will need further validation with additional disease models and human data. Other poorly understood metabolic relationships that require further investigations include the crosstalk between tumor immune cell and adipocyte metabolism and the tumor microenvironment and cancer cell survival.

Concluding Remarks

Comorbidities should be considered throughout the cancer research process. The advent of COVID-19 infections is reinforcing the notion that diseases other than cancer decrease cancer survival [1]. Many chronic diseases have adverse effects on cancer outcomes, leading to excessive mortality among cancer patients. These deaths are preventable with treatment, diet and lifestyle changes, increased physical activity, and psychosocial support, among other intervention strategies that target these chronic diseases. Comorbidities influence tumor biology through various mechanisms (Figure 1). It is important that we understand these biology-based mechanisms to improve prevention and intervention as part of cancer care, specifically for patient populations that are at increased risk of multiple comorbidities like the elderly. There are many unanswered questions (see Outstanding Questions). For the most part, we have a limited understanding of how comorbidities affect therapy response in the elderly, underserved, and race/ethnic minorities, in part because clinical trials exclude those with common comorbidities (Figure 2). Certain comorbidities have been studied more extensively, such as obesity and diabetes, but others have been neglected, like liver and renal diseases, metabolic syndrome, neurological disorders, and autoimmune diseases. Lack of suitable animal cancer models to study comorbidities may have contributed to the dearth in mechanistic research. Likewise, funding opportunities for this research may not exist. We rarely have patient data that support our findings from *in-vitro* and animal studies. One reason could be the difficulty of obtaining biospecimens from cancer patients with known comorbidities. Comorbidities may affect younger subjects differently than elderly subjects and women differently than men. Age and gender are modifiers of health and disease and their influence remains to be understudied and underestimated in clinical research and medical practice, including the effects of comorbidities on cancer outcomes. Our understanding of tumor immune cell function and cancer metabolism is still in an evolving state. For example, we do not know how comorbidities affect tumor immune response through metabolic changes in immune cells. There are some data for adipocytes, endothelial cells, and tumor-associated fibroblasts, but they are generally sparse. Hence, more mechanistic research is needed into comorbidity-induced cancer progression and mortality for increased awareness among clinicians, and to develop novel prevention and intervention strategies.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Glossary

Hyperglycemia

Refers to a high blood sugar (glucose) level. Usually occurs when the body does not produce insulin or when insulin signaling is compromised. High blood sugar is an indicator of diabetes.

PD-1 blockade

PD-1 or Programmed cell death protein 1 is a cell surface protein that regulates the immune system's response to cells in the human body by down-regulating the immune response. PD-1 inhibitors cause a PD-1 signaling blockade and can activate the immune system to attack tumors.

Gut microflora/microbiome

The microbiome represents all microbes - bacteria, fungi, protozoa and viruses - that live on or inside the human body. In our body, the gut microflora/microbiome is the largest entity. The microbiome consists of microbes that are both helpful and potentially harmful.

Dysbiosis

Indicates a deleterious microbial imbalance in the body, usually an imbalance of the gut microflora.

Epithelial to mesenchymal transition (EMT)

EMT is a reversible cellular program by which epithelial cells lose their cell polarity and cell-cell adhesion, and gain migratory and invasive properties to become mesenchymal type cells like fibroblasts. The mesenchymal differentiation allows cancer cells to become more metastatic.

Reactive oxygen species

A type of unstable molecules that contain oxygen and easily react with other molecules in cells. They may cause damage to cell components such as DNA, RNA, lipids, and proteins and can cause mutations and cell death. However, reactive oxygen species can also participate in signal transduction.

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Box 1.**Comorbidities can alter the gut microbiome**

The gut microbiome affects human health. A dysbiosis can increase cancer risk. Comorbidities may confer their cancer risk through effects on the gut microbiome. There is evidence that a dysbiosis can be a cause of cancer [41, 42]. Diabetes can lead to depletion of beneficial butyrate-producing taxa in the human gut microbiota [43], whereas hyperglycemia can cause intestinal barrier dysfunction and enteric infections [44]. A high-fat diet can promote intestinal carcinogenesis in *K-ras* mutant mice by causing a gut microbiota dysbiosis, driven by a decrease in Paneth-cell-mediated antimicrobial host defense and compromised dendritic cell recruitment [45]. The gut microbiome also promotes obesity-associated liver cancer by inducing deoxycholic and lipoteichoic acid synthesis and a senescence-associated secretory phenotype in hepatic stellate cells with increases in inflammatory and tumor promoting factors and prostaglandin E2-mediated suppression of antitumor immunity [46, 47]. Recently, it was shown that the gastrointestinal microbiome influences efficacy of PD-1-based cancer immunotherapy in mice and patients, largely due to certain commensal species that define the clinical response [48]. The study indicated that an existing dysbiosis as a comorbidity may negatively affect immunotherapy efficacy. Furthermore, comorbidities may contribute to persistent gene expression alterations and epigenome remodeling that predispose to cancer or promote disease progression [49–51]. An obesity-associated microbiome was found to reprogram the intestinal epigenome as well, leading to persistent alterations that may promote carcinogenesis [52].

Outstanding Questions

Can we validate findings from animal models of comorbidities with patient data? Can we show that comorbidities in patients associate with a distinct tumor biology that is related to findings from experimental research? How do comorbidities relate to the development of an aggressive tumor biology in patients?

There is good evidence that obesity, diabetes, chronic infections, and stress exposures, can alter the tumor immune environment and gut microflora, but most other cancer-associated comorbidities have not been studied in this context. How do other cancer-associated comorbidities, like the metabolic syndrome, chronic cardiovascular disease, liver and renal disease, autoimmune diseases, or depression, affect the tumor immune environment and the gut microflora in cancer patients?

What is the effect of comorbidities on cancer therapy response in patient groups at high risk of multiple comorbidities, such as the elderly, underserved, and race/ethnic minorities?

Is the effect of comorbidities on tumor biology dependent on patients' age or gender?

How do comorbidities affect the metabolism and function of all tumor-associated cells, including cancer cells, immune and endothelial cells, adipocytes, and tumor-associated fibroblasts?

Highlights

The burden of comorbid chronic diseases is increasing as populations get older and is affecting underserved populations more so than the affluent

Comorbidities and cancer have common risk factors, but comorbidities and medications to treat them can directly or indirectly influence tumor biology

Comorbidities have adverse effects on cancer prevention and outcomes and lead to an excessive cancer mortality that is preventable

They exert their effects on tumor biology by altering the tumor microenvironment, cancer metabolism, and the gut microflora, and by affecting the cancer therapy response

They promote cancer progression by mechanisms including local and systemic inflammation, changes to the tumor immune environment, mesenchymal differentiation of cancer cells, and formation of a pro-metastatic niche at distant organ sites

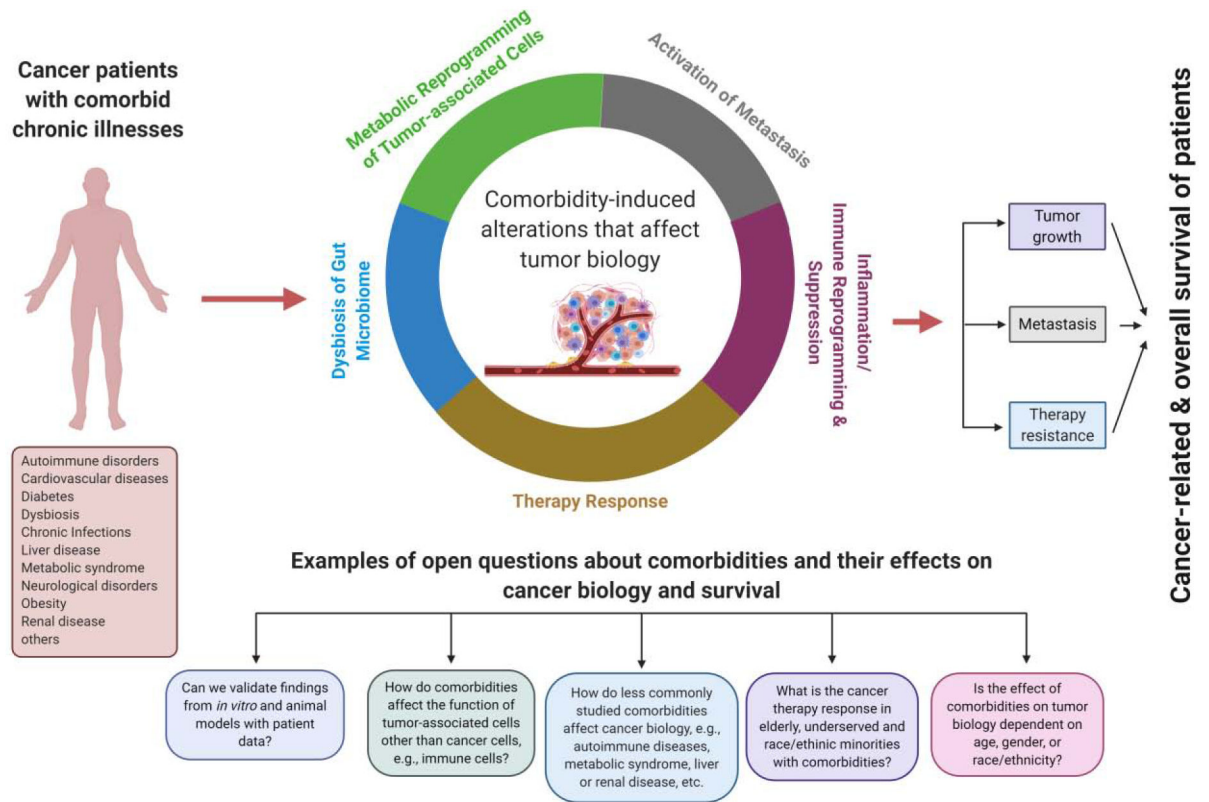


Figure 1: Current knowledge and open questions about cancer-related comorbidities and their effects on cancer biology and survival

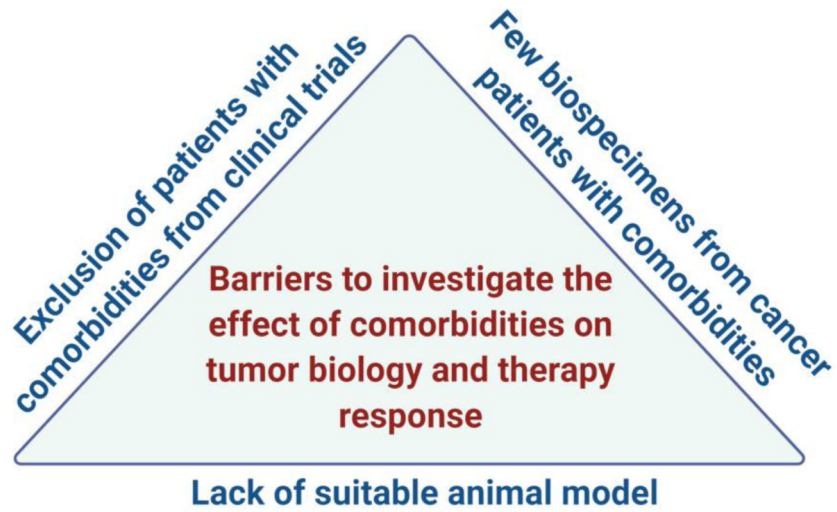


Figure 2:
Barriers to studying the adverse effects of cancer-related comorbidities