Overview

Hot and cold tumor are crucial in assessing the efficacy of existing cancer immunotherapies. A hot tumor is rich in tumor immune microenvironment (TME), PD-L1 overexpression, genomic instability. Example of hot tumors include melanoma and lung cancer. In contrast, non-T- cell inflamed cancers, such as Triple Negative Breast Cancer (TNBC), prostate or pancreas cancers fall into the category of "cold tumors". Simultaneously, tumor-associated macrophages (TAMs), T/B regulatory cells (T/Bregs), myeloid-derived suppressor cells (MDSCs) have been identified as contributor to the formation of a specific TME in "cold tumors". Factors affecting tumor virulence include immune checkpoints (PD-1, PD-L1, CTL4, TIM-3, and LAG-3), tumor infiltrating lymphocytes (TIL), tertiary lymphoid tissue (TLS), microsatellite status (MSI), tumor mutational burden (TMB) and gene mutations (for ex.TP53, BRCA1).

Immune checkpoint inhibitors (ICIs) have shown success in improving the survival of cancer patient. ICI-mediated antitumor responses depend on the infiltration of T cells that identify and eliminate cancer cells. Therefore, ICIs are less efficacious in "cold tumors" which are characterized by the lack of T-cell infiltration.

Specific Aims

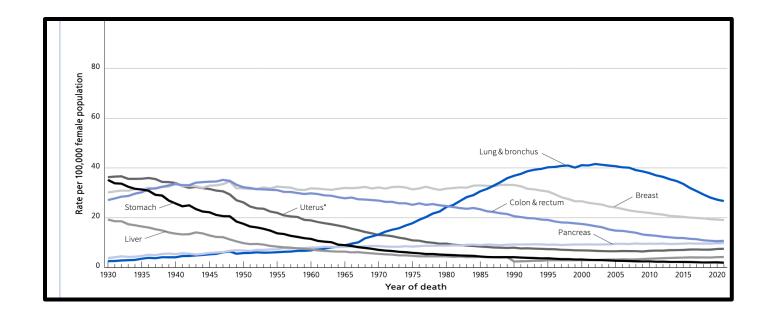
The aims of this project are to analyze the current landscape of strategies designed in "warming up" cold tumors to immune checkpoint inhibitors (ICIs); to introduce an innovative approach utilizing nanoparticles, viral vectors, or polymeric particles as potential solutions. We will also sketch a clinical trial with appropriate criteria selection of patients to validate the established solution.

Significance

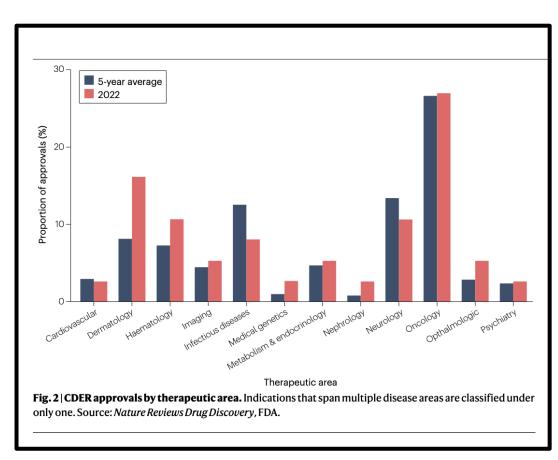
In last 10 years, the FDA has approved an increasing number of Immune Checkpoint Inhibitors (ICIs) following successful clinical trials. These treatments have significantly enhanced long-term survival rates for metastatic patients and prolong progression-free survival for those in early stages of the disease; yet some patients fail to respond to these strategies; in addition, these drugs are ineffective for certain tissue or tumor types. For 2024, in the United States there will be around 2 million new cancer diagnoses, with an estimated of 9,620 cases affecting children and 5,290 adolescents. This year is also expected to bring about approximately 611,720 cancer-related deaths, including 1,040 children and 550 adolescents.

According to the National Cancer Institute, cancer-related healthcare expenses in the U.S. reached \$208.9 billion in 2020. Since their peak in 1991, there has been a 33% decline in the rates of most common cancers, including lung, colorectal, breast and prostates, The trend has been attributed to a combination of factors, such as reduced smoking rates, advanced in therapies like ICIs, and the development of improved diagnostic and prognostic biomarkers. Nonetheless, cancer incidence rates have increased in breast, uterine, melanoma and prostate cancers [1].

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In 2022, the FDA approved 37 new drugs, with ICIs leading the charge. Among these approvals, there were new checkpoint inhibitors like LAG-3, as well novel antibodies and targeted gene therapies, such as those targeting KRAS G12C mutation in NSCLC [2]. However, while these therapies hold great promise, patients can still experience autoimmune side effects causing lifethreatening complications affecting



various organs (please refer to side effects reported by American Cancer Society).

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Cold tumors are characterized by a deficiency in T cells, and in the absence of T cells, there are no checkpoint inhibitors to activate. Several key factors contribute to tis shortage of T cells, including:

- Insufficient tumor antigens: cold tumors exhibit a scarcity of tumor-specific antigens, hindering the immune system's ability to recognize effectively.
- Inadequate antigen processing and presentation machinery (APM)
- Low Tumor Mutational Burden (TMB): cold tumors typically have fewer mutations and a lower load of neoantigens, making them less recognizable to the immune system.
- Downregulation of MHC-I molecules: reduction in class I MHC molecules limits the
 presentation of tumor antigens, diminishing the immune system's ability to target the
 tumor cells. In addition, tumor cells can inhibit DC phagocytosis and escape the
 immune system.
- A variety of oncogenic pathways, such as WNT/Bet-catenin, KRAS or MYC; could be activated to regulate the immune response. Factors such as low levels of proinflammatory cytokines can contribute to immunosuppression. By contrast, stromal cells (CAFs) can redirect CTLs toward the cancer stroma preventing them from entering the tumor. Elevated expression of CXCL8 has been linked to a decrease of T cell presence and an increase in neutrophils within the tumor microenvironment (TME).
- Inadequate vasculature, resulting from ineffective aggregation of vascular endothelial adhesion cells, reduce T-cell trafficking to the tumor. Vascular endothelial growth factor (VEGF) triggers the formation of new blood vessels, which in turn affects the movement of T cells toward the tumor. Furthermore, factors such as hypoxia driven by the transcription factors HIF1, acidosis, and necrosis, contribute to the recruitment of immunosuppressive cells within the TME. These conditions also induce angiogenesis through molecules like CCL28 and VEGF), further impacting T-cell trafficking and influencing the presence of inflammatory molecules like CD39 and CD73.
- Immunosuppressive cells and factors: within the cancer stroma, cancer-associated fibroblasts (CAFs) play a pivotal role as they produce an extracellular matrix that acts as a physical barrier. CALFs also release CXCL12 which T-cell response and produce TGFBeta which limits the proliferation of CD4+ T cells. TGF-Beta achieves this by inhibiting production of IL-2 and promoting the conversion of naïve CD4 + into Treg.
- Elevated glycolysis activity and the subsequent accumulation of lactate are associated with a negative correlation to the infiltration of CD8+ cells, often indicated by high expression of LDH-A. Similarly, heightened cholesterol levels have been observed to downregulate MHC-I levels resulting in a decreased infiltration of CTLs.

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1] "2024-cancer-facts-and-figures-acs.pdf".

[2] "d41573-023-00001-3.pdf".

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