

A Detailed Review on Cancers: It's Physical and Biochemical Basis

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Abstract

This thesis presents a comprehensive review of cancer, with a particular focus on its physical and biochemical basis. Cancer, a leading cause of mortality worldwide, is characterized by the uncontrolled growth and spread of abnormal cells. Understanding the fundamental differences between cancerous and normal cells is essential for developing effective diagnostic tools and therapeutic strategies.

The review begins by exploring the physical anomalies of cancer cells, including their irregular shape, size and structural organization. These cells often exhibit altered cell membranes, disrupted cytoskeletal architecture and irregularities in cell adhesion and communication, which contribute to their invasive and metastatic potential.

The biochemical basis of cancer is examined in detail, with an emphasis on the dysregulation of key metabolic pathways. Cancer cells undergo profound metabolic reprogramming, often referred to as the Warburg effect, where they preferentially utilize glycolysis over oxidative phosphorylation, even in the presence of oxygen. This metabolic shift supports the rapid proliferation of cancer cells and their survival in hypoxic conditions. Additionally, cancer cells produce abnormal proteins that contribute to tumor progression, such as mutant p53 and fail to produce critical tumor suppressor proteins like RB1, leading to unchecked cell division and evasion of apoptosis.

Furthermore, the thesis reviews the role of oncogenes and tumor suppressor genes in cancer development, highlighting how mutations and epigenetic alterations drive the transformation of normal cells into malignant ones. The complex interplay between these genetic changes and the tumor microenvironment is also discussed, emphasizing the importance of signaling pathways, immune evasion and angiogenesis in cancer progression.

This detailed review synthesizes current research and insights from authoritative sources, including peer-reviewed journals and foundational medical texts, to provide a thorough understanding of the physical and biochemical underpinnings of cancer. The findings underscore the importance of continued research in these areas to advance the diagnosis, treatment and prevention of cancer.

Keywords: Cancer; Immunology; Warburg effect; Metabolic pathways; Cytoskeletal architecture

Introduction

Cancer is a complex group of diseases marked by the uncontrolled division and growth of abnormal cells in the body. The term "cancer" encompasses over 100 different types of malignancies, each of which originates from various cells and tissues. It represents a significant public health challenge, being one of the leading causes of morbidity and mortality worldwide.

Background and definition

At its core, cancer arises when normal cellular processes are disrupted, leading to unregulated cell growth. Under normal circumstances, cells in the human body grow, divide and die in a controlled manner. However, when genetic mutations occur within a cell, these regulatory processes can be altered or completely bypassed, causing cells to proliferate uncontrollably. This unchecked growth can lead to the formation of a mass of tissue known as a tumor.

Tumors can be classified as benign or malignant. Benign tumors are generally localized, grow slowly and do not invade other tissues. They are usually not life-threatening. In contrast, malignant tumors are aggressive, can invade nearby tissues and have the potential to spread to distant parts of the body through a process known as metastasis [1].

Origins and etiology

The transformation of a normal cell into a cancerous one is typically driven by genetic mutations, which can be inherited or acquired. These mutations often affect genes involved in cell cycle regulation, apoptosis (programmed cell death), DNA repair and other critical cellular functions. Several factors contribute to the initiation and progression of cancer, including:

Genetic pre-disposition: Some individuals inherit mutations that significantly increase their risk of developing specific cancers. For instance, mutations in the *BRCA1* and *BRCA2* genes are strongly associated with an increased risk of breast and ovarian cancers.

Environmental factors: Exposure to carcinogens such as tobacco smoke, Ultraviolet (UV) radiation, certain chemicals and infectious agents like Human Papillomavirus (HPV) can induce mutations in DNA, leading to cancer. For example, smoking is the leading cause of lung cancer, while UV exposure is a major risk factor for skin cancers.

Lifestyle factors: Diet, physical inactivity, alcohol consumption and obesity have been linked to an increased risk of various cancers. For instance, diets high in processed meats and low in fruits and vegetables are associated with colorectal cancer.

Age: The risk of developing cancer increases with age, as the accumulation of genetic mutations over time can lead to malignant transformations in cells [2].

Literature Review

Cellular and molecular basis

The process by which normal cells transform into cancerous cells is known as carcinogenesis, which typically involves multiple steps:

Initiation: This stage involves the occurrence of genetic mutations within a cell, often due to exposure to carcinogens or other damaging agents.

Promotion: In this phase, the initiated cells are stimulated to divide and proliferate. This process is often driven by factors such as hormones, chronic inflammation or additional genetic changes.

Progression: The final stage involves further genetic alterations that confer aggressive characteristics to the cancer cells, such as the ability to invade tissues and metastasize to other parts of the body.

On a molecular level, cancer cells exhibit several distinct features that differentiate them from normal cells. These include the ability to sustain proliferative signaling, evade growth suppressors, resist cell death, enable replicative immortality, induce angiogenesis (formation of new blood vessels) and activate invasion and metastasis [3]. Collectively, these traits are often referred to as the "hallmarks of cancer."

Cancer is a multifaceted disease with a complex interplay of genetic, environmental and lifestyle factors. Understanding the underlying mechanisms of cancer development is crucial for developing effective prevention strategies and therapeutic interventions. As research advances, a deeper comprehension of cancer biology continues to improve the prospects for early detection, targeted treatment and ultimately, the prevention of this devastating disease.

Physical properties

Size and shape

Normal cells: Generally, normal cells have a consistent, regular shape and size. Their dimensions and forms are tightly regulated by the body, maintaining tissue architecture.

Cancerous cells: Cancer cells often display significant variations in size and shape, a phenomenon known as pleomorphic. They can be larger or smaller than normal cells, with irregular, distorted shapes. This variation is one of the hallmarks of malignancy.

Nuclear-cytoplasmic ratio

Normal cells: The Nucleus-to-Cytoplasm (N:C) ratio in normal cells is typically low. The nucleus is proportionately smaller compared to the cytoplasm.

Cancerous cells: Cancer cells often have a higher N:C ratio, meaning the nucleus is disproportionately large compared to the cytoplasm. This is due to an increased amount of genetic material and a higher rate of cellular replication.

Cell membrane

Normal cells: The cell membrane in normal cells is smooth, with regular patterns of surface markers (like antigens) that maintain cell communication and function.

Cancerous cells: In cancerous cells, the cell membrane may have irregularities, including an increased number of surface receptors, loss of cell adhesion molecules and altered glycoproteins. These changes can facilitate unregulated growth and invasion into surrounding tissues.

Cellular architecture

Normal cells: Normal cells are typically organized in a manner that reflects their function within tissues. For instance, epithelial cells line up in a regular pattern.

Cancerous cells: Cancerous cells often lose this organized structure, leading to anaplasia (loss of differentiation) and dysplasia (disordered growth). This disorganization allows cancer cells to grow uncontrollably.

Internal composition

Genetic material (DNA and chromosomes)

Normal cells: Normal cells have a stable genetic material, with a regular number of chromosomes that follow the diploid state.

Cancerous cells: Cancer cells frequently exhibit genetic instability, including mutations, chromosomal aberrations (like translocations, duplications or deletions) and aneuploidy (an abnormal number of chromosomes). These genetic changes contribute to the uncontrolled growth and survival of cancer cells.

Mitochondria

Normal cells: Mitochondria in normal cells function efficiently, providing energy primarily through oxidative phosphorylation.

Cancerous cells: In cancerous cells, mitochondrial function is often altered, with a shift towards glycolysis even in the presence of oxygen, known as the Warburg effect. This metabolic reprogramming supports rapid cell division and growth [4].

Cytoskeleton

Normal cells: The cytoskeleton, which includes microfilaments, intermediate filaments and microtubules, maintains the cell's shape and aids in intracellular transport.

Cancerous cells: Cancer cells often have altered cytoskeletal structures, which can contribute to their ability to invade and metastasize. Changes in actin filaments and microtubules are particularly important in enabling cell motility and invasion.

Cell cycle regulation

Normal cells: Normal cells are tightly regulated by various checkpoints during the cell cycle (G₁, S, G₂ and M phases). This regulation ensures controlled growth and division.

Cancerous cells: Cancer cells often bypass these regulatory checkpoints due to mutations in key genes such as p53, RB and APC. This allows them to proliferate uncontrollably, contributing to tumor growth.

Apoptosis mechanism

Normal cells: Apoptosis or programmed cell death, is a controlled process in normal cells that eliminates damaged or unneeded cells.

Cancerous cells: Cancer cells often evade apoptosis by altering the expression of pro-apoptotic and anti-apoptotic proteins, such as Bcl-2, Bax and caspases. This evasion of apoptosis is a critical feature of cancer cells, allowing them to survive beyond their normal lifespan.

Angiogenesis

Normal cells: Normal cells do not typically induce the formation of new blood vessels, except during wound healing or in specific physiological conditions.

Cancerous cells: Cancer cells can promote angiogenesis (the formation of new blood vessels) by secreting factors like Vascular Endothelial Growth Factor (VEGF). This supports their growth by providing necessary nutrients and oxygen and also facilitates metastasis.

Metabolism

Normal cells: Normal cells primarily utilize oxidative phosphorylation in mitochondria to produce ATP, with glycolysis being a secondary pathway.

Cancerous cells: Cancer cells often rely more heavily on glycolysis even under aerobic conditions (aerobic glycolysis or Warburg effect). This metabolic shift supports rapid cell proliferation and survival in hypoxic environments often found within tumors.

Molecular and biochemical differences

Oncogenes and tumor suppressors

Normal cells: In normal cells, proto-oncogenes (which promote cell growth) and tumor suppressor genes (which inhibit cell growth) are in balance, ensuring regulated cell proliferation and survival.

Cancerous cells: In cancer cells, mutations often activate oncogenes (e.g., Ras, Myc) and inactivate tumor suppressor

genes (e.g., TP53, BRCA1/2), leading to uncontrolled cell division and tumorigenesis.

Signal transduction pathways

Normal cells: Signal transduction pathways in normal cells are precisely regulated, ensuring that growth signals are balanced with inhibitory signals.

Cancerous cells: Cancer cells often have dysregulated signaling pathways, such as the PI3K/AKT/mTOR pathway, which drives proliferation, survival and growth independent of external growth factors.

Epigenetic changes

Normal cells: Epigenetic modifications in normal cells, like DNA methylation and histone modification, are tightly controlled and help regulate gene expression.

Cancerous cells: In cancer cells, abnormal epigenetic modifications can lead to the silencing of tumor suppressor genes and activation of oncogenes. These changes are often reversible, making them a target for therapeutic interventions.

Discussion

The transition from normal cells to cancerous cells involves profound changes in both physical properties and internal composition [5,6]. These alterations allow cancer cells to evade normal regulatory mechanisms, grow uncontrollably, invade surrounding tissues and eventually metastasize.

Structural anomalies cell wall/cell membrane

Normal cells: The cell membrane in normal cells is composed of a lipid bilayer with embedded proteins, maintaining structural integrity and regulating material passage. The membrane proteins include receptors, channels and adhesion molecules that play roles in signal transduction and cell communication.

Cancerous cells: The cell membrane of cancer cells often shows significant structural alterations:

Increased fluidity: Cancer cell membranes tend to be more fluid due to altered lipid composition, which aids in metastatic behavior. This is partly because of changes in the ratio of cholesterol to phospholipids and alterations in fatty acid saturation levels.

Loss of polarity: Normal epithelial cells have apical-basal polarity that is crucial for their function. Cancer cells often lose this polarity, contributing to their disorganized structure and ability to invade other tissues.

Altered glycocalyx: The glycocalyx (a carbohydrate-rich zone on the cell surface) in cancer cells is often altered, with changes in glycoproteins and glycolipids. This can enhance the cell's ability to evade immune detection and adhere to other cells or extracellular matrix, aiding in metastasis.

Receptors

Normal cells

Receptor function: Normal cells have a variety of receptors that regulate cell growth, differentiation and apoptosis. These include growth factor receptors (e.g., EGFR), hormone receptors (e.g., estrogen receptors) and immune cell receptors.

Balanced signaling: In normal cells, receptor activation is tightly regulated. Ligand binding leads to downstream signaling that promotes or inhibits cellular functions as needed.

Cancerous cells

Overexpression of receptors: Cancer cells often exhibit overexpression of growth factor receptors (e.g., HER2 in breast cancer) and other receptors that drive proliferation. This leads to enhanced and uncontrolled signaling for growth and division.

Mutated receptors: Mutations in receptors, such as the Epidermal Growth Factor Receptor (EGFR) in non-small cell lung cancer, can lead to constant activation even in the absence of ligands, driving the malignant phenotype [7].

Loss of tumor suppressor receptors: Tumor suppressor receptors that inhibit growth (e.g., TGF- β receptors) are often downregulated or mutated in cancer cells, leading to a loss of inhibitory control on cell proliferation.

Altered immune receptors: Cancer cells can also express altered or increased immune checkpoint receptors (e.g., PD-L1) to evade immune surveillance, allowing the tumor to grow unchecked.

Material exchange mechanisms

Transport mechanisms in normal cells

Selective permeability: Normal cell membranes are selectively permeable, allowing controlled passage of ions, nutrients and waste products *via* passive (diffusion, facilitated diffusion) and active (pumps, carriers) transport mechanisms.

Ion channels and pumps: Sodium-potassium pumps, calcium channels and other ion channels maintain ionic gradients essential for cellular function and signaling.

Transport mechanisms in cancer cells

Increased nutrient uptake: Cancer cells often upregulate transporters for glucose (e.g., GLUT1) and amino acids to meet the increased metabolic demands of rapidly dividing cells. This is part of the metabolic reprogramming seen in many cancers.

Acidification of microenvironment: Cancer cells frequently exhibit increased expression of proton pumps and ion exchangers (e.g., Na⁺/H⁺ exchangers), contributing to an acidic extracellular environment. This acidity promotes invasion and inhibits immune cell function.

Drug resistance mechanisms: Cancer cells can overexpress efflux pumps (e.g., P-glycoprotein) that remove chemotherapeutic agents from the cell, contributing to multidrug resistance.

Endocytosis and exocytosis mechanisms

Endocytosis in normal cells

Regulated uptake: Normal cells use endocytosis to internalize nutrients, hormones and other molecules. This process is tightly regulated and occurs *via* clathrin-mediated endocytosis, caveolae-mediated endocytosis or macropinocytosis, depending on the cargo.

Receptor-mediated endocytosis: Ligand binding to surface receptors often triggers their internalization *via* clathrin-coated pits, allowing cells to regulate receptor levels and signaling.

Endocytosis in cancer cells

Enhanced endocytosis: Cancer cells often show increased rates of endocytosis to sustain their high metabolic needs and to internalize more nutrients and growth factors.

Altered pathways: The endocytosis pathways can be altered in cancer cells. For instance, macropinocytosis is often upregulated in cancers like pancreatic cancer, allowing cells to uptake large amounts of extracellular nutrients.

Exosome production: Cancer cells release more exosomes (*via* exocytosis), which are small vesicles that carry proteins, lipids and nucleic acids. These exosomes can modulate the tumor microenvironment, promote metastasis and suppress immune responses.

Exocytosis in normal cells

Controlled exocytosis: Exocytosis in normal cells involves the secretion of cellular products like enzymes, hormones and neurotransmitters. This process is controlled and occurs through vesicle docking and fusion with the plasma membrane.

Vesicle formation: Vesicles containing cellular products are formed in the Golgi apparatus and transported to the plasma membrane for exocytosis.

Exocytosis in cancer cells

Increased secretion: Cancer cells often have enhanced exocytosis, particularly of proteases and Matrix Metalloproteinases (MMPs), which degrade the extracellular matrix and facilitate invasion and metastasis.

Immune modulation: The exocytosis of immune-modulatory molecules, such as cytokines and chemokines, by cancer cells can alter the immune environment to favor tumor growth and inhibit anti-tumor immunity.

Drug resistance: Cancer cells can use exocytosis to secrete chemotherapeutic agents, contributing to drug resistance. This exocytosis is often mediated by upregulated vesicle transport proteins.

Specific examples in major common cancers

Breast cancer

HER2 overexpression: In certain breast cancers, there is an overexpression of HER2 receptors, leading to increased signaling for cell growth and division. This is often targeted by drugs like trastuzumab [8].

Exosome release: Breast cancer cells release exosomes containing microRNAs that can suppress immune function and promote metastasis.

Lung cancer

EGFR mutations: Non-small cell lung cancers frequently have mutations in the EGFR receptor, leading to constant activation and uncontrolled proliferation.

Altered glycolysis: These cells often exhibit increased expression of glucose transporters, enhancing glycolysis and energy production to support rapid growth.

Colorectal cancer

Wnt pathway activation: In colorectal cancer, mutations in the APC gene lead to constitutive activation of the Wnt signaling pathway, promoting cell proliferation.

Altered cell junctions: There is a loss of E-cadherin in colorectal cancer cells, which disrupts cell-cell adhesion and contributes to metastasis [9].

Pancreatic cancer

KRAS mutations: Mutations in the KRAS gene lead to hyperactivation of downstream signaling pathways that promote cancer cell survival and proliferation.

Macropinocytosis: Pancreatic cancer cells often rely on macropinocytosis to uptake extracellular protein, which is broken down into amino acids to fuel their growth.

Conclusion

Cancer cells exhibit profound alterations in their structure, cell membrane, receptors, material exchange mechanisms and endocytosis/exocytosis processes compared to normal cells. These changes are integral to the cancer cell's ability to grow uncontrollably, evade immune surveillance, invade surrounding tissues and metastasize to distant organs. Understanding these differences provides critical insights into cancer biology and highlights potential targets for therapeutic interventions.

Cancer cells exhibit several significant biochemical anomalies compared to normal cells, particularly in terms of metabolism, energy sources, energy utilization mechanisms and the production or lack of specific proteins. Below is a detailed exploration of these differences:

Metabolism and energy sources

Glycolysis and the Warburg effect: Cancer cells predominantly rely on aerobic glycolysis for energy production, even in the presence of oxygen, a phenomenon known as the Warburg effect. This contrasts with normal cells, which typically rely on oxidative phosphorylation in the mitochondria under aerobic conditions. The preference for glycolysis in cancer cells allows them to produce energy more rapidly, though less efficiently, to support rapid cell division and growth. The byproduct of this process, lactate, accumulates in the tumor microenvironment, leading to acidification, which can promote tumor progression and immune evasion.

Lipid metabolism: Alterations in lipid metabolism are also a hallmark of cancer cells. They show increased lipogenesis (fatty acid synthesis) to support membrane formation and produce signaling molecules essential for cell survival and proliferation. For example, the enzyme Stearoyl-CoA Desaturase 1 (SCD1) is upregulated in many cancers, facilitating the conversion of saturated fatty acids to monounsaturated fatty acids, which are crucial for maintaining membrane fluidity and supporting tumor growth and resistance to therapy.

Amino acid metabolism: Cancer cells have a heightened demand for certain amino acids, particularly glutamine, which serves as a carbon and nitrogen source for biosynthesis and energy production. Glutamine is used in the Tricarboxylic Acid (TCA) cycle to produce ATP and generate antioxidants like glutathione, protecting cancer cells from oxidative stress. In some cancers, glutamine is so crucial that it becomes a limiting factor for survival and proliferation. Cancer cells also upregulate transporters for amino acids like serine, glycine and branched-chain amino acids to support rapid growth.

Energy utilization mechanisms

Oxidative Phosphorylation (OXPHOS) vs. Glycolysis: While normal cells rely on OXPHOS in the mitochondria for efficient ATP production, cancer cells shift towards glycolysis, which, despite being less efficient, provides metabolic intermediates necessary for biosynthesis. This metabolic shift is partly driven by mutations in oncogenes and tumor suppressor genes, such as MYC and p53, which reprogram the energy metabolism of cancer cells to favor growth over efficiency.

Lipid and amino acid metabolism: Beyond glycolysis, cancer cells also modify lipid and amino acid metabolism to meet their energy and biosynthesis needs. Enhanced fatty acid oxidation (FAO) provides an additional energy source, especially under nutrient stress. Similarly, the metabolism of certain amino acids like glutamine supports the anabolic needs of proliferating cancer cells, contributing to the synthesis of nucleotides, proteins and lipids.

Abnormal protein production

Oncogenes and tumor suppressors: Cancer cells often produce abnormal proteins due to mutations in oncogenes (e.g., KRAS, BRAF) and tumor suppressors (e.g., p53). These proteins can drive uncontrolled cell proliferation, resist apoptosis (programmed cell death) and enable metastasis. For instance, mutant p53 proteins not only lose their tumor-suppressive functions but can also gain new oncogenic properties, contributing to tumor progression.

Lactate Dehydrogenase (LDH): Cancer cells often overproduce LDH, which facilitates the conversion of pyruvate to lactate in the glycolytic pathway. Elevated levels of LDH are associated with poor prognosis in various cancers, as they promote an acidic microenvironment that supports invasion, metastasis and immune escape.

Deficiencies in essential proteins

p53: In many cancers, p53, a critical tumor suppressor protein, is either mutated or inactivated. The absence or malfunction of p53 disrupts the cell's ability to undergo apoptosis in response to DNA damage, allowing cancer cells to survive and proliferate despite genomic instability. This loss is a significant contributor to cancer development and resistance to therapy.

BRCA1/2: These proteins are involved in DNA repair through homologous recombination. In cancers such as breast and ovarian cancers, mutations in BRCA1/2 impair DNA repair, leading to the accumulation of genetic mutations that drive tumorigenesis. The loss of BRCA1/2 function also renders cells more sensitive to certain DNA-damaging agents, a vulnerability that can be exploited therapeutically.

The failure of the immune system to detect and destroy cancer cells is a complex process that involves several mechanisms by which cancer cells evade immune surveillance. Here's a detailed explanation of why this happens and the strategies being explored to trigger an effective immune response against cancer:

Immune evasion mechanisms of cancer cells

Cancer cells develop several strategies to escape the immune system, allowing them to grow and spread unchecked:

Reduced antigen presentation: Cancer cells can lose or downregulate the expression of tumor antigens or Major Histocompatibility Complex (MHC) molecules on their surface. This makes them less visible to T cells, which rely on recognizing these antigens to initiate an immune response.

Immunosuppressive Tumor Microenvironment (TME): The TME is often rich in immunosuppressive cells like regulatory T cells (Tregs), Myeloid-Derived Suppressor Cells (MDSCs) and Tumor-Associated Macrophages (TAMs). These cells secrete cytokines such as IL-10 and TGF- β , which inhibit the activation and function of cytotoxic T cells and Natural Killer (NK) cells.

Immune checkpoint activation: Tumors can express immune checkpoint proteins like PD-L1, which bind to PD-1 receptors on T cells. This interaction inhibits T cell activation, allowing the cancer cells to evade immune destruction. Similarly, other checkpoints like CTLA-4, LAG-3 and TIGIT also play roles in dampening the immune response immunotherapy).

Secretion of immunosuppressive factors: Cancer cells can secrete various factors, including Indoleamine 2,3-Dioxygenase (IDO), which depletes tryptophan and suppresses T cell function. They can also release extracellular vesicles containing proteins or RNA that modulate immune responses in favor of tumor survival.

Strategies to trigger an immune response against cancer

To counteract these evasion mechanisms, several strategies are being developed and implemented:

Immune checkpoint inhibitors: These drugs block inhibitory signals (e.g., PD-1/PD-L1, CTLA-4) that prevent T cells from attacking cancer cells. By inhibiting these checkpoints, the

immune system can be reactivated to target and kill cancer cells more effectively.

Adoptive Cell Therapy (ACT): This involves extracting T cells from a patient's tumor, expanding or genetically modifying them to enhance their cancer-fighting abilities and reinfusing them into the patient. This approach boosts the number of T cells that can target the tumor.

Modulation of the tumor microenvironment: Strategies that alter the immunosuppressive TME are being explored, such as inhibiting Treg or MDSC function or repolarizing TAMs to support anti-tumor immunity. Drugs targeting these cells and their secreted factors can help shift the balance towards a more immune-supportive environment.

Oncolytic viruses: These are engineered viruses that selectively infect and kill cancer cells while also stimulating an immune response against the tumor. They can also be used to deliver immune-activating genes directly to the tumor site.

Challenges and future directions: Despite these advancements, challenges remain in overcoming immune resistance. Tumors are heterogeneous and can develop resistance to immunotherapy over time. Combining different immunotherapeutic strategies or using them alongside traditional therapies like chemotherapy and radiation is an ongoing area of research. Personalized medicine, which tailors treatment based on the genetic and immunologic profile of an individual's tumor, is also showing promise in enhancing the effectiveness of cancer immunotherapy.

There are several promising, yet underexplored, approaches that could potentially harness the immune system's response against cancer cells by targeting their distinct physical and biochemical properties compared to normal cells. These emerging strategies focus on exploiting the unique mechanical and biochemical features of the tumor microenvironment and cancer cells to provoke an immune response. Here's a detailed exploration of these possibilities:

Mechanotransduction in immune cells: Mechanotransduction, the process by which cells convert mechanical stimuli into biochemical signals, is a critical aspect of immune cell function. Immune cells such as T-cells, Natural Killer (NK) cells and macrophages are influenced by the mechanical properties of their environment, including the stiffness and composition of the Extracellular Matrix (ECM) around tumors. Cancer cells often alter their ECM, making it stiffer, which can affect immune cell behavior. By designing materials or therapies that mimic or modulate these mechanical cues, it might be possible to enhance the activation and targeting capabilities of immune cells against tumors. For instance, the activation of mechanosensitive ion channels like TRPV4 in immune cells could be targeted to boost immune responses against cancer.

Biophysical properties and immune targeting: Cancer cells exhibit unique biophysical properties, such as altered cell membrane stiffness, increased cell surface area due to membrane protrusions (e.g., lamellipodia and invadopodia) and enhanced interaction with the ECM. These features are critical for cancer cell migration and invasion. Targeting these

biophysical differences can provoke an immune response. For example, nanoparticles or biomimetic materials designed to recognize and bind to the altered mechanical properties of cancer cells could be used to deliver immune-modulating agents directly to tumors, enhancing the immune system's ability to detect and destroy cancer cells.

Immune cell mechanosensation: NK cells and macrophages are particularly sensitive to the mechanical properties of their surroundings. Research suggests that these cells could be engineered or stimulated to respond more aggressively to the altered mechanical cues present in the tumor microenvironment. For example, enhancing the mechanosensitive pathways within these immune cells could improve their ability to infiltrate tumors and kill cancer cells. This could be achieved by developing drugs or therapies that modulate these pathways, potentially leading to new forms of cancer immunotherapy.

Targeting integrin-mediated adhesion: Integrins are transmembrane receptors that mediate the adhesion between cells and the ECM, playing a crucial role in cell migration and signal transduction. Cancer cells often exhibit increased integrin activity, which supports their invasive behavior. By developing therapies that specifically target the integrins involved in cancer cell-ECM interactions, it may be possible to disrupt these processes and enhance immune recognition and response to the tumor. This approach could be particularly effective in combination with existing immunotherapies, providing a new avenue for cancer treatment immunotherapies.

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