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# Mechanisms by which obesity regulates inflammation and anti-tumor immunity in cancer

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### ABSTRACT

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Obesity is associated with an increased risk for 13 different cancers. The increased risk for cancer in obesity is mediated by obesity-associated changes in the immune system. Obesity has distinct effects on different types of inflammation that are tied to tumorigenesis. For example, obesity promotes chronic inflammation in adipose tissue that is tumor-promoting in peripheral tissues. Conversely, obesity inhibits acute inflammation that rejects tumors. Obesity therefore promotes cancer by differentially regulating chronic versus acute inflammation. Given that obesity is chronic, the initial inflammation in adipose tissue will lead to systemic inflammation that could induce compensatory anti-inflammatory reactions in peripheral tissues to suppress chronic inflammation. The overall effect of obesity in peripheral tissues is therefore dependent on the duration and severity of obesity. Adipose tissue cellularity is different ant different anatomical sites throughout the body. Consequently, the sensitivity of adipose tissue to obesity is dependent on the anatomical location of the adipose depot. For example, obesity induces more inflammation in visceral than subcutaneous adipose tissue. Based on these studies, the mechanisms by which obesity promotes tumorigenesis are multifactorial and immune cell type-specific. The objective of our paper is to discuss the cellular mechanisms by which obesity promotes tumorigenesis by regulating distinct types of inflammation in adipose tissue and the tumor microenvironment.

### 1. Introduction

Obesity is a widespread chronic disease [1-10]. Obesity is a risk factor for cancer [11–20]. Specifically, obesity increases the risk for 13 different cancers [21]. The prevalence of worldwide obesity has approximately doubled since 1980 such that nearly one-third of the adult world population is now overweight or obese [4]. It is projected that nearly 60 % of the world's population will be overweight or obese by the year 2030 [4]. It is estimated that one in five men and women will develop cancer worldwide and that one in three adults will be overweight or obese, indicating that the majority of adults who will have cancer will also have obesity [4,12]. The global increase in obesity is hypothesized to contribute to the increasing incidence of early-onset cancers, such as colon cancer [22]. The increase in breast cancer incidence also coincides with the surge in obesity [23]. Obesity is positively associated with breast cancer risk in postmenopausal women and inversely correlated with breast cancer risk in premenopausal women [23]. The difference in obesity association with breast cancer based on menopausal age is in part due to estrogen synthesis by breast adipose tissue [23]. Obesity is also linked with increased risk and progression of breast cancer that is negative for the ER, progesterone receptor, and HER2 [24,25]. The increasing incidence of uterine cancer is hypothesized to be in part due to obesity [26–28]. The link between obesity and uterine cancer is especially strong considering obesity increases the risk for this hormone-responsive tumor by 7-fold [21]. Cancer is increased in obesity by several mechanisms, including the promotion of chronic inflammation, and the inhibition of acute inflammation that rejects tumors. Herein, we will review the cellular mechanisms that are involved in obesity-mediated chronic inflammation and suppression of acute inflammation that mediates anti-tumor immunity.

# 2. Adipokines

### 2.1. Cytokine effects of leptin

The term adipokine refers to a large group of bioactive molecules released from an adipocyte. Many adipokines have roles in cancer, and

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| Abbreviations |   | LepR              | Leptin Receptor   |
|---------------|---|-------------------|---|
|               |   | MCP1              | Monocyte Chemoattractant Protein 1                      |
| APC           | Antigen-Presenting Cells                              | MDSCs             | Myeloid Derived Suppressor Cells                        |
| ATP           | Adenosine Triphosphate                                | MHC               | Major Histocompatibility Complex                        |
| BC            | breast cancer   | MMP2              | Matrix Metalloproteinases 2                             |
| BRCA          | Breast Cancer Associated Gene                         | MMP9              | Matrix Metalloproteinases 9                             |
| CAA           | Cancer Associated Adipocytes                          | mTOR              | Mammalian Target of Rapamycin                           |
| CD4           | Cluster of Differentiation 4                          | NASH              | nonalcoholic steatohepatitis                            |
| CD8           | Cluster of Differentiation 8                          | NK                | Natural Killer cells                                    |
| CRC           | colorectal carcinoma                                  | PAI1 –            | Plasminogen Activator Inhibitor 1                       |
| CTRP3         | Adipokine C1q/Tumor Necrosis Factor Related Protein 3 | PD-1              | Programmed Cell Death 1                                 |
| CTRP9         | Adipokine C1q/Tumor Necrosis Factor Related Protein 9 | PD-L1             | Program Death Ligand 1                                  |
| DNA           | Deoxyribonucleic Acid                                 | PHD3              | Prolyl Hydroxylase 3                                    |
| ERK           | Extracellular Signal Related Kinase                   | RANTES            | Regulated on Activation Normal T Expressed and Secreted |
| FGFB21        | Fibroblast Growth Factor 21                           | RBP4              | Retinol Binding Protein 4                               |
| GLP-1 –       | Glucagon Like Peptide –1                              | RNA               | Ribonucleic acid  |
| HCC           | hepatocellular carcinoma                              | SCC               | squamous cell carcinoma                                 |
| HUVEC         | Human Umbilical Vein Endothelial Cells                | STAT              | Signal Transducer and Activator of Transcription 3      |
| IGFBP2        | Insulin-Like Growth Factor Binding Protein 2          | TAMs              | Tumor-Associated Macrophages                            |
| IGF1          | Insulin-Like Growth Factor 1                          | T <sub>H</sub> 1  | T helper cell 1   |
| IL-1β         | Interleukin 1 Beta                                    | T <sub>H</sub> 17 | T helper cell 17  |
| IL-2 –        | Interleukin 2   | TME               | Tumor Microenvironment                                  |
| IL-6 –        | Interleukin-6   | TNBC              | Triple Negative Breast Cancer                           |
| IL-10         | Interleukin 10  | $TNF\alpha -$     | Tumor Necrosis Factor Alpha                             |
| IL-17 –       | Interleukin 17  | Tregs             | T regulatory cells                                      |
| JAK           | Janus Kinase  | VEGF              | Vascular Endothelial Growth Factor                      |
|               |   |                   |   |

in this review, we have summarized their effects on cancer in Table 1. The adipokine leptin has been intensely studied, and we have reviewed its signaling mechanisms in more detail. The crystal structure of leptin is homologous to the long-chain helical cytokine family that induces IL-6 [29]. Leptin signals through its cell surface Jak-STAT linked receptor [30]. The binding of leptin to the leptin receptor (LepR), induces LepR dimerization and JAK2-mediated phosphorylation of specific residues in the intracellular domain of the LepR [30]. The phosphorylation of JAK2 sites in the LepR mediates the induction of STAT3 and ERK signaling in response to leptin [30]. Leptin acts on hypothalamic neurons to induce a satiety effect [31]. The increase in fat mass in obesity is correlated with an increase in serum leptin concentrations. The serum level of leptin in lean humans is 5 ng/mL and in obese humans, the levels of leptin can reach 100 ng/mL [32]. Overproduction of leptin leads to leptin resistance in obesity [31]. Clinically, leptin resistance causes overeating and obesity [31]. Leptin resistance may be cell-type specific. For example, obese levels of leptin are associated with the progression of breast cancer [33]. Leptin when applied to breast cancer cells in cell culture induces a concentration-dependent increase in signaling with a maximal effect observed in response to an obese concentration (100 ng/mL) of leptin [34]. Breast cancer cells also express leptin and the LepR [35]. The concentration of leptin in some tumors therefore might be higher than the concentration of leptin in blood. Most immune cell types in innate and adaptive T cell immunity express the leptin receptor (LepR) [36]. Broadly, the role of leptin in immunity is to boost inflammatory reactions. For instance, leptin augments  $TNF\alpha$  and  $IL-1\beta$  production by human macrophages [37]. Leptin may also promote the initiation of cancer. For example, leptin stimulates an increase in DNA damage in normal breast epithelial cells obtained from women with a mutation in the BRCA gene [38]. The mechanism by which leptin promotes DNA damage is unknown.

### 3. Obesity-associated inflammation

Localized acute inflammation in adipose tissue is a beneficial mechanism that protects against determinantal deposition of lipids in

peripheral tissues such as the liver [165]. In obesity, however, the chronic overfilling of adipocytes with triglycerides causes chronic low-grade inflammation in adipose tissue that leads to low-grade systemic inflammation [166,167]. The cytokines that emerge from adipose tissue modulate inflammatory pathways in peripheral organs [166]. Given that adipose tissue is in all tissues, chronic inflammation can occur in all tissues in response to obesity [166]. However, the severity of obesity-induced adipose tissue inflammation is dependent on the anatomical location of the adipose tissue in the body [168-170]. For example, in obesity, visceral fat has more inflammation than subcutaneous fat [171-176]. In humans, white adipocytes make up 98 % of total fat mass and the remaining adipocytes are brown adipocytes [177]. White adipocytes store and release energy and initiate inflammation in response to obesity. Brown adipocytes regulate thermogenesis and are less prone to obesity-associated inflammation [178-180]. In humans, body fat distribution is also linked with differences in risk for cardiovascular disease. The accumulation of abdominal fat promotes the risk of cardiovascular disease and insulin resistance. Conversely, gluteal-femoral subcutaneous fat protects against these obesity-associated diseases [181-188]. Visceral adipose tissue contains a greater percentage of hypertrophic adipocytes, and immune cells, and thus is more sensitive to obesity than subcutaneous adipose tissue. This difference could in part explain why the accumulation of visceral fat is more strongly associated with metabolic disease compared to an increase in the amount of subcutaneous adipose tissue [188]. Obesity also has distinct effects on different types of inflammation. For example, obesity promotes chronic inflammation in adipose tissue, yet inhibits acute inflammation in the tumor microenvironment (TME). In this review, we will discuss the cellular mechanisms by which obesity promotes low-grade inflammation, and inhibits acute inflammation in the TME.

# 3.1. Inflammation promotes cancer risk

There are several examples by which chronic inflammation promotes cancer and these studies support the hypothesis that obesity, by

### Table 1

Summary table of the biomolecules secreted from adipose tissue, the effect obesity has on their secretion, and their role in cancer pathogenesis. Abbreviations: CRC- colorectal carcinoma, HCC- hepatocellular carcinoma, BC- breast cancer, NASH- nonalcoholic steatohepatitis, SCC- squamous cell carcinoma, TNBC- Triple Negative Breast Cancer, PD-L1- Program Death Ligand 1, mTOR-Mammalian Target of Rapamycin, HUVEC- human umbilical vein endothelial cells, RNA- Ribonucleic acid, DNA- Deoxyribonucleic Acid.

| Adipokine Name   | Obesity<br>Effect | Effect on Cancer  | Reference  |
|--|-------------------|---|------------|
| Leptin   | <b>↑</b>          | <ul> <li>Increase cell<br/>proliferation in breast<br/>cancer</li> <li>Increase cell invasion<br/>in CRC</li> </ul>   | [39–44]    |
|  |                   | <ul> <li>promote secretion of<br/>exosomes</li> </ul>   |            |
| Adiponectin  | ŧ                 | <ul> <li>Triggers cell death in<br/>BC</li> <li>Suppress cell</li> </ul>  | [40,45–47] |
|  |                   | migration in prostate cancer cells  |            |
| Visfatin   | <b>↑</b>          | <ul> <li>Promotes TNBC cell<br/>stemness</li> <li>Anoikis resistance in<br/>ovarian cancer</li> </ul>   | [48–51]    |
| Fetuin-A   | <b>†</b>          | unknown   | [52,53]    |
| Plasminogen Activator<br>Inhibitor 1 (PAI-1)                           | <b>†</b>          | <ul> <li>Marker of poor<br/>prognosis in breast<br/>and ovarian cancer</li> <li>inhibits cell migration<br/>in pancreatic cancer,<br/>glioma, and<br/>melanoma</li> </ul> | [54–59]    |
| Resistin   | <b>1</b>          | <ul> <li>Prostate cancer cell<br/>proliferation</li> </ul>  | [60–63]    |
| Omentin-1  | ŧ                 | <ul> <li>Inhibits proliferation<br/>and promotes<br/>apoptosis in CRC</li> <li>promotes expression<br/>of tumor suppressive<br/>microRNA in CRC</li> </ul>                | [64–67]    |
| Lipocalin-2  | <b>†</b>          | <ul> <li>Inhibits ferroptosis in<br/>CRC</li> <li>Promotes TME in<br/>hepatocellular<br/>carcinoma HCC</li> </ul>   | [68–71]    |
| Asprosin   | <b>†</b>          | <ul> <li>promote progress to<br/>invasive SCC</li> </ul>  | [72–74]    |
| Neuregulin 4   | ŧ                 | <ul> <li>serves as a hormonal<br/>checkpoint to prevent<br/>progression of NASH-<br/>HCC</li> </ul>   | [75–78]    |
| Vaspin   | <b>1</b>          | <ul> <li>Increase proliferation,<br/>migration and<br/>invasion of TNBC</li> </ul>  | [79–81]    |
| Insulin-like Growth<br>Factor Binding<br>Protein-2 (IGFBP-2)           | ŧ                 | <ul> <li>promotes metastasis<br/>of pancreatic cancer<br/>and chemotherapy<br/>resistance</li> </ul>  | [82–84]    |
| Fibroblast Growth<br>Factor 21 (FGF21)                                 | <b>†</b>          | unknown   | [85,86]    |
| Retinol Binding Protein<br>4 (RBP4)                                    | <b>†</b>          | <ul> <li>increases ovarian<br/>cancer migration</li> </ul>  | [87–89]    |
| Adipokine C1q/Tumor<br>Necrosis Factor<br>Related Protein 3<br>(CTRP3) | Controversial     | - promotes<br>proliferation of<br>osteosarcoma cells  | [90–93]    |
| Adipokine C1q/Tumor<br>Necrosis Factor<br>Related Protein 9<br>(CTRP9) | <b>↑</b>          | unknown   | [94,95]    |

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Table 1 (continued)

| Adipokine Name  | Obesity<br>Effect | Effect on Cancer  | Reference         |
|---|-------------------|---|-------------------|
| Irisin  | <b>↑</b>          | <ul> <li>inhibit tumor<br/>development and<br/>induce apoptosis in</li> </ul>   | [96–100]          |
| Apelin  | t                 | prostate cancer<br>- promotes prostate<br>cancer progression<br>- increased tumor size,<br>stage, and poor<br>prognostic measure in<br>BC   | [101–104]         |
| Angiopoietin like<br>protein 2  | <b>+</b>          | <ul> <li>increase breast cancer<br/>metastasis</li> <li>chemotherapy<br/>registrance in CPC</li> </ul>  | [105–108]         |
| Chemerin  | <b>†</b>          | <ul> <li>promotes endothelial<br/>angiogenesis and<br/>migration of HUVEC<br/>cells</li> </ul>  | [109–111]         |
| Progranulin   | <b>†</b>          | <ul> <li>upregulates PDL-1 in<br/>breast cancer</li> <li>promotes growth,<br/>migration, and<br/>invasion of CRC</li> </ul>   | [112–115]         |
| Monocyte<br>chemoattractant<br>Protein 1 (MCP1/<br>CCL2)                            | <b>†</b>          | <ul> <li>poor prognostic<br/>marker in multiple<br/>cancers including<br/>breast, lung, and<br/>prostate cancer</li> <li>induce tamoxifen<br/>resistance</li> </ul>   | [112,<br>116–121] |
| Matrix<br>Metalloproteinases 2<br>(MMP2)  | <b>↑</b>          | <ul> <li>promotes invasion<br/>and migration in<br/>bladder and prostate<br/>cancer</li> </ul>  | [122–125]         |
| Matrix<br>Metalloproteinases 9<br>(MMP9)  | <b>↑</b>          | <ul> <li>promotes invasion<br/>and migration in<br/>bladder and prostate<br/>capeer</li> </ul>  | [122–125]         |
| Activin A   | -                 | <ul> <li>increased invasion,<br/>migration and<br/>metastasis in prostate,<br/>BC, and squamous cell<br/>carcinoma SCC</li> <li>induces apoptosis in<br/>HCC</li> <li>suppresses<br/>angiogenesis and<br/>tumor growth in<br/>gastric cancer</li> </ul> | [126–132]         |
| Regulated on<br>Activation, Normal T<br>expressed and<br>Secreted (RANTES/<br>CCL5) | <b>†</b>          | <ul> <li>promotes immune cell<br/>infiltration of tumors</li> <li>glioblastoma<br/>chemotherapy<br/>resistance</li> </ul>   | [133–136]         |
| Insulin Like Growth<br>Factor-1 (IGF-1)   | Controversial     | <ul> <li>increased tumor<br/>development,<br/>angiogenesis, and<br/>metastasis in CRC</li> </ul>  | [137–140]         |
| Lysophosphatidic Acid   | Ť                 | <ul> <li>increase proliferation<br/>and migration of<br/>prostate cancer cells</li> <li>suppress autophagy<br/>through activation of<br/>mTOR in prostate<br/>cancer</li> </ul>   | [141–144]         |
| Vascular Endothelial<br>Growth Factor<br>(VEGF)                                     | <b>†</b>          | <ul> <li>mediator of<br/>angiogenesis in<br/>cancer</li> <li>suppress antitumor<br/>immune activity</li> </ul>  | [145–148]         |
| Interleukin-1 beta (IL-<br>1β)  | t                 | <ul> <li>marker of high-grade<br/>dysplasia or cancer</li> <li>promote<br/>carcinogenesis</li> </ul>  | [149–152]         |

(continued on next page)

### Table 1 (continued)

| Adipokine Name  | Obesity<br>Effect | Effect on Cancer  | Reference                      |
|---|-------------------|---|--------------------------------|
| Interleukin-6 (IL-6)  | •                 | <ul> <li>promotes resistance<br/>to anti-PDL-1<br/>immunotherapy</li> <li>expansion of cancer<br/>stem cell population<br/>in BC</li> </ul>   | [82,<br>153–156]               |
| Tumor Necrosis Factor<br>alpha (TNF α)  | <b>↑</b>          | <ul> <li>breast cancer<br/>progression and<br/>metastasis</li> <li>promotes epithelial to<br/>mesenchymal<br/>transition</li> <li>promotes suppression<br/>of T-regulatory cells</li> </ul> | [157–160]                      |
| Extracellular Vesicles<br>- siRNA<br>- microRNA<br>- DNA<br>- Lipids<br>- Sphingolipids<br>- Phosphatidylserine | Unknown           | <ul> <li>promote cell to cell<br/>communication in<br/>breast cancer</li> <li>influence tumor<br/>microenvironment</li> </ul>   | [161,162],<br>[44,163,<br>164] |

promoting low-grade inflammation, promotes tumorigenesis. For example, the presence of gut pathogens promotes inflammatory bowel disease, which increases the risk of colon cancer [189]. Human papillomavirus induces cervicitis and increases the risk of cervical cancer [189]. The risk for hepatocellular carcinoma is increased by hepatitis, which is caused by the hepatic B/C virus [189]. The Epstein-Barr virus induces mononucleosis and is a risk factor for Burkitt's lymphoma [189]. The hypothesis that chronic inflammation in obesity promotes cancer is supported by the link between cancer and other instances of inflammation, including chronic inflammation.

# 3.2. Obesity-associated mechanisms of chronic inflammation in adipose tissue

The inflammatory environment in adipose tissue is highly responsive to obesity. With obesity, the inflammatory milieu in adipose tissue moves from anti-inflammatory to pro-inflammatory. This regulation is mediated by changes in the numbers and activity of anti- and proinflammatory immune cells in adipose tissue in response to obesity. For example, regulatory T cells (Tregs) inhibit inflammation by releasing the anti-inflammatory cytokine, interleukin 10 (IL-10) [190]. During murine and human obesity, there is a reduction in Tregs and IL-10 in adipose tissue [190-193]. Macrophages are also responsive to obesity [194,195]. There is increased infiltration and phenotypic switching to proinflammatory M1-stage macrophages in adipose tissue in obesity [194-196]. M1-stage macrophages release tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ), interleukin 6 (IL-6), and IL-1 $\beta$  [197]. The recruitment of macrophages is due to chemokines released from necrotic adipocytes in obese adipose tissue [198]. The inflammatory activity of M1-macrophages in adipose tissue is further increased by leptin which is produced in greater amounts by adipocytes in obesity [37]. Obesity increases the number of inflammatory T helper 1 (T<sub>H</sub>1), and T<sub>H</sub>17 cells in part by upregulating major histocompatibility complex class II on adipocytes in adipose tissue [192]. The primary cytokine from  $T_{\rm H}1$  cells is interferon-gamma (INF- $\gamma$ ). INF- $\gamma$  is also the primary cytokine that downregulates the number of Tregs, which stimulates more inflammation in adipose tissue in obesity [191].  $T_{\rm H}1$  cells also release TNF and IL-2 cytokines [199]. Cytotoxic CD8 T cells are also responsive to obesity. The induction and expansion of adipose tissue-resident CD8 T cells initiate the recruitment of proinflammatory M1-macrophages to adipose tissue in obesity [200,201]. T<sub>H</sub>17 are inflammatory T cells that are increased in adipose tissue in obesity [202,203]. IL-17 is the proinflammatory cytokine that is primarily released by  $T_{\rm H}17$  cells [204].

Functionally, IL-17 activates neutrophils, which are the first immune cells to infiltrate adipose tissue in obesity [205]. IL-17 also inhibits adipocyte differentiation and therefore the increase in IL-17 in obesity disrupts adipose tissue [203]. Clinically, IL-17 could be linked to autoimmunity in obese humans [206]. Fig. 1 summarizes the mechanisms of obesity-associated inflammation in adipose tissue. Based on these studies, the accumulation of inflamed adipose tissue causes systemic inflammation that increases cancer risk in obesity.

### 4. Breast adipose tissue inflammation

Clinically, breast adipose tissue inflammation is measured by the number of tissue slides that are positive for the presence of macrophage crown-like structures surrounding dying adipocytes [207-210]. The density of crowns per tissue section is also reported [207-210]. The macrophages in crowns surrounding dying adipocytes are immunostained with an anti-CD68 antibody, which is a pan-marker of macrophages [207–210]. The presence of macrophage crown-like structures is a good readout of inflammation, given that the infiltration of macrophages is the first and most robust inflammatory response in adipose tissue in obesity [211,212]. Clinical studies show obesity is significantly associated with increased abundance of macrophage crowns in breast adipose tissue in women with breast cancer [207-210]. Obesity-associated systemic increases in leptin, IL-6, and triglycerides are also positively correlated with breast adipose tissue inflammation in women who underwent mastectomy for breast cancer treatment [209]. In women, breast adipose tissue inflammation is positively associated with faster relapse of metastatic breast disease [209]. The progression of benign breast disease to breast cancer is also significantly associated with breast adipose tissue inflammation [207]. The presence of crowns in breast adipose tissue was significantly higher in women with breast cancer compared with women who did not have cancer [210]. Women who were carriers of a BRCA mutation did not show increased breast adipose tissue inflammation [210]. Breast adipose tissue inflammation



Fig. 1. Adipose tissue inflammation in obesity.

Adipose tissue is highly responsive to obesity-induced inflammation. The increase in MCP-1, leptin, cytotoxic CD8 T cells, and activated neutrophils recruit and polarize proinflammatory M1-macrophages in adipose tissue in obesity. There is increased development of proinflammatory TH1 and TH17 cells, which by releasing INF<sub>q</sub> and IL-17 inhibit Tregs and stimulate neutrophils, respectively. Obesity-associated adipose tissue inflammation increases circulating TNF- $\alpha$ , IL-6, and leptin. MCP-1 = Monocyte Chemoattractant Protein 1, TNF $\alpha$  = Tumor necrosis factor-alpha, IL-6 = Interleukin-6, M1 = M1-stage Macrophage, IL-7 = Interleukin-17, TH17 = T helper 17, TH1 = T helper 1, Treg = Regulatory T cells, INF<sub>q</sub> = Interferon-gamma, IL-10, Interleukin-10.

is not associated with breast tumor grade [208]. However, breast adipose tissue crowns occurred more frequently in women with luminal-B-like breast tumors [210]. The presence of breast adipose tissue crowns was associated with increased lymphovascular disease for triple-negative breast cancer (TNBC) [210]. The mechanisms that link luminal-B-like breast tumors and TNBC invasiveness with increased breast adipose tissue inflammation in women have not been published. Studies have asked if the presence of breast adipose tissue crowns is associated with breast cancer progression. The abundance of macrophage crowns in breast adipose tissue is not associated with disease-free progression or overall survival of breast cancer patients [208,210]. The role of race has also been investigated, and the results show that the number of breast adipose tissue crowns in African-American women was not significantly different than the number of breast adipose tissue crowns in White women with breast cancer [208]. The role of the tumor has been assessed and the results show the abundance of macrophage crowns in breast adipose tissue is higher in obese women with breast cancer compared with obese women who do have cancer [210]. This finding indicates the tumor secretes proinflammatory cytokines that act on breast adipose tissue. Combining multiple biomarkers for breast adipose tissue inflammation could be a more accurate predictor of cancer progression than just assessing the number of macrophage crowns in breast adipose tissue. A future study that combines multiple markers of breast adipose tissue inflammation could identify an improved strategy that might be significantly associated with disease-free survival and overall survival for breast cancer patients.

### 5. Anti-tumor immunity

The recognition and killing of cancer cells by T cells is a seven-step process [213,214]. The first step starts with the release of cancer-specific antigens by the tumor. The cancer antigens are detected and processed by antigen-presenting cells (APCs). Cancer-primed APCs activate T cells in the lymph node. Cancer-primed cytotoxic T cells (CD8<sup>+</sup> T cells) traffic in the bloodstream to the tumor. Cancer-activated CD8<sup>+</sup> T cells infiltrate the tumor and react with cancer cells that present antigens via the major histocompatibility complex (MHC). The final step is the killing of cancer cells by CD8<sup>+</sup> T cells. Natural killer (NK) cells also contribute to the killing of cancer cells through an MHC-independent mechanism [215]. Tumors evade anti-tumor immunity by downregulating the expression of MHC and upregulating the expression of PD-L1 on cancer cells [216-218]. PD-L1 expressed on cancer cells binds to PD-1 expressed on T cells and this association between PD-L1 and PD-1 induces signaling in T cells that inhibits T cell effector activity [219-221]. Tregs and myeloid-derived suppressor cells (MDSCs) are immunosuppressive and they promote cancer by inhibiting anti-tumor immunity [222,223]. Consequently, increased numbers of Tregs and MDSCs in the tumor microenvironment are associated with poor cancer prognosis [224,225].

Upon entering a tumor, a subset of effector-like  $CD8^+$  T cells differentiate into tissue-resident memory T cells [226]. Tissue-resident T cell markers are CD69 or CD103, which retain T cells in tissue through integrin interactions [226]. In peripheral tissues, memory resident T cells release granzyme B in response to antigen stimulation, which can eliminate pathogen-infected cells [226]. Tissue-resident T cells also release the pro-inflammatory cytokines INF $\gamma$ , TNF $\alpha$ , and IL-2, which recruit additional inflammatory cells to the site of infection [226]. The release of granzyme B and proinflammatory cytokines by tumor resident T cells suppress tumor growth [226]. Tumor resident T cells are heterogeneous, with subsets expressing several T cell exhaustion and activation markers [226]. Clinically, increased number of resident CD8<sup>+</sup> T cells in tumors is associated with good cancer prognosis [226]. In the sections below, we will review how obesity inhibits anti-tumor immunity by acting on T cells, NK cells, and MDSCs.

### 6. The roles of obesity in anti-tumor immunity

### 6.1. Upregulation of PD-1 on T cells in obesity

Upon their recruitment to the TME, CD8 T-cells become exhausted [227]. T cells that become exhausted are initially active, but their persistent overstimulation in the TME leads to their gradual loss of effector activity and their inability to kill cancer cells [227]. The upregulation of PD-1 on T cells is a marker of T cell exhaustion [228]. The PD-1 that is expressed on T cells binds to PD-L1, which is expressed on the surface of cancer cells. The binding of PD-1 to PD-L1 induces signaling in T cells that block T cell effector activity [221]. The hallmark of an exhausted CD8 T cell in the TME is the inability of the PD-1-expressing CD8 T cell to kill a cancer cell [221]. Consequently, obesity by increasing the expression of PD-1 on T cells facilitates T cell inactivation in the TME and this favors tumorigenesis [229]. Mechanistically, leptin acts on T cells to induce phosphorylated STAT3 signaling that upregulates the expression of PD-1 [229] (Fig. 2). Higher levels of leptin in obesity, therefore, drive the inactivation of CD4<sup>+</sup> helper T cells, and CD8<sup>+</sup> cytotoxic T cells [229]. The induction of PD-1 on T cells in response to obesity was confirmed in mice, primates, and humans and therefore this mechanism could promote tumorigenesis in obese patients [229]. The induction of PD-1 on CD8<sup>+</sup> T cells in response to obesity was associated with reduced CD8<sup>+</sup> T cell effector activity and reduced release of cytokines (IL-2 and interferon-gamma) [229]. Thus, the induction of PD-1 on T cells in obesity is linked to the functional suppression of CD8<sup>+</sup> T cells. RNA-seq experiments coupled with pathway analysis showed that CD8<sup>+</sup> T cells from obese mice exhibited gene expression changes significantly associated with metabolic alterations, and T cell hyporesponsiveness, such as T cell anergy [229]. Thus, the observed reduction in CD8<sup>+</sup> T cell activity in obesity is due to





Increased leptin in obesity acts on cytotoxic CD8 T cells to induce signaling that promotes the phosphorylation of STAT3 (Tyr705) which in turn activates the STAT3 response element within the enhancer of the PD-1 gene which leads to increased PD-1 expression by T cells. The binding of T cell-expressed PD-1 to cancer cell-expressed PD-L1 induces signaling in CD8 T cells that inactivates the effector function of T cells, leading to an exhausted T cell state. P-STAT3 = Phosphorylated-Signal Transducer And Activator Of Transcription 3 (Tyr705), PD-1 = Programmed cell death protein 1, PD-L1 = Programmed Death-1 (PD-1) Lizand 1.

multiple changes in the T cell that inhibit T cell effector activity towards cancer cells [229]. Preclinically, the effect of PD-1 inhibitors on tumor growth in obese compared with lean mice was investigated [229]. Mechanistically, PD-1 antibodies, by binding to PD-1 on T cells, block PD-1 binding to PD-L1 on cancer cells [221]. PD-1 blocking drugs by preventing the binding of T cell PD-1 to cancer cell PD-L1 disinhibits the T cell, and the now active CD8<sup>+</sup> T cell kills cancer cells [221]. Of interest, is that inhibiting PD-1 with PD-1 antibody, inhibited the growth of melanoma and mammary tumors in obese mice more than it inhibited the growth of these tumors in lean mice [229]. A similar response has been reported in humans with melanoma, such that obese patients are more responsive to PD-1 inhibitors than lean patients [229]. From an immunity standpoint, the PD-1 inhibitor not only inhibited the growth of tumors in obese mice, but it also increased the number of activated CD8<sup>+</sup> T cells, and M1-phase macrophages, and reduced the number of immunosuppressive MDSCs in the TME and peripheral tissues [230]. These findings show that blocking PD-1 activity in obese mice with cancer, restored several aspects of anti-tumor immunity and that this was correlated with a reduction in the growth of tumors in obese mice, and the anti-tumor effect was stronger in obese, than lean mice [229]. The greater efficacy of blocking PD-1 translates to humans, given that the efficacy of PD-1 inhibitors in obese humans with cancer is better than the efficacy of PD-1 inhibitors in lean humans with cancer [231]. This human response has been shown for melanoma, lung cancer, and renal cancer in human obesity [231]. Mechanistically, it is hypothesized that CD8<sup>+</sup> T cells are more responsive to PD-1 blockers in obesity because CD8<sup>+</sup> T cells are more suppressed by PD-1 signaling in obese compared with lean mice or humans [231].

# 6.2. Obesity inhibits the acquisition of free fatty acids by T cells in the TME

There is competition between T cells and cancer cells for nutrients in the TME [232]. Cancer cells out-compete T cells for nutrients by upregulating the expression of nutrient transporters and increasing their utilization of nutrients [232]. This, in turn, establishes a gradient by which nutrients flow from the tumor interstitium into cancer cells, instead of moving into T cells [232]. The reduced flow of nutrients into T cells prevents the expansion and effector activity of T cells in the TME and this compromises anti-tumor immunity and the tumor evades the immune system. Prior reports have focused on the reduced flow of glucose and amino acids into T cells due to enhanced uptake of these nutrients by cancer cells [232]. In obesity, however, the gradient is free fatty acids, such that these lipids flow into and are rapidly utilized by cancer cells. In contrast, the movement of free fatty acids into CD8<sup>+</sup> T cells in the TME is suppressed [233]. The reduced flow of lipids into  $CD8^+$  T cells is associated with reduced  $CD8^+$  T cell infiltration in tumors and poor CD8<sup>+</sup> T cell effector activity in obese mice compared with lean mice with tumors [233]. Interestingly, this process of free fatty acids preferentially being moved into cancer cells at the expense of CD8<sup>+</sup> T cells is induced by the downregulation of prolyl hydroxylase 3 (PHD3) in cancer cells in obesity [233] (Fig. 3). In lean mice, PHD3 is highly expressed in cancer cells, and it blocks the movement of free fatty acids into the mitochondria; therefore, lipids are not readily utilized by cancer cells in lean mice, and this allows lipids to distribute into CD8<sup>+</sup> T cells within the TME, and this is associated with improved anti-tumor immunity in lean mice [233]. However, in obesity, PHD3 is downregulated in cancer cells [233] (Fig. 3). In response to PHD3 downregulation, free fatty acids flow into the mitochondria of cancer cells in obesity, and this increased utilization of free fatty acids by the mitochondria for fatty acid oxidation drives a gradient by which free fatty acids in the tumor interstitium are transported into cancer cells, not CD8<sup>+</sup> T cells, and this is associated with inhibited CD8<sup>+</sup> T cell effector activity in the tumors of obese mice [233]. Interestingly, genetically preventing the downregulation of PHD3 in cancer cells in obesity inhibits lipid uptake by cancer cells, restores CD8<sup>+</sup> T cell effector activity,



Fig. 3. T cell inactivation in obesity.

There is a competition between T cells and cancer cells for limiting amounts of FFAs in the tumor interstitium. In lean mice, reduced utilization of FFAs by cancer cells due to upregulation of PHD3 promotes the flow of FFAs into T cells. In obese mice, reduced PHD3 promotes FFA utilization by cancer cells, which diverts the flow of FFAs away from T cells. PHD3 = Prolyl hydroxylase 3, FFA = Free Fatty Acids, FAO = Fatty Acid Oxidation.

and inhibits the growth of tumors in obese mice [233]. Clinically, analysis of the Cancer Genome Atlas showed the levels of PHD3 were negatively correlated with BMI in the context of cancer [233]. These findings provide the premise for a future study investigating what factor (s) in obesity reduce PHD3 expression in cancer cells. Hypothetically, preventing obesity-stimulated downregulation of PHD3 in cancer cells would inhibit tumor growth by restoring anti-tumor immunity. In addition to altering fatty acid and lipid metabolism, obesity decreases amino acid metabolism in tumor-resident CD8<sup>+</sup> T cells in MC38 colon tumors [234]. This is secondary to reduced activity of the amino acid transporter, SLC7A5, on tumor resident T cells in response to obesity [234]. Conversely, leucine uptake by breast cancer cells is stimulated by obesity-associated breast adipokines [235]. Increased leucine uptake by breast cancer cells is mediated by increased SLC7A5 activity in response to obesity-associated adipokines [235]. These findings show that tumor-associated T-cell activity and cellular metabolism are interconnected and dysfunctional in obesity.

# 6.3. Natural killer (NK) cells

NK cells are inhibited in obesity [236,237]. In the context of cancer, NK cells synapse with a tumor cell, and polarize their cytolytic granules to release a payload of proteases and hydrolases onto cancer cells, which kills the cancer cell [238]. In obesity, NK cells synapse with tumor cells, yet they fail to polarize and, therefore, fail to release their cytolytic granules onto cancer cells, and the cancer cell is not lysed [236]. This defect in the polarization of lytic granules is linked to a defect in glycolysis and oxidative metabolism in NK cells in obesity [236]. The defect in glycolysis and oxidative metabolism is due to the accumulation of triglyceride storage droplets in the cytoplasm of NK cells in murine and human obesity [236,237]. In obesity, NK cells accumulate cytoplasmic lipid droplets via a transcriptional mechanism that promotes lipid absorption and synthesis [237]. In NK cells, excess lipid accumulation in the cytoplasm inhibits the activity of mTOR and destabilizes P300 and cMYC protein [237]. The mTOR pathway, P300, and cMYC drive metabolism in NK cells [236,237]. The metabolic defect in NK cells in obesity is therefore linked to the loss of mTOR, P300, and cMYC [236, 237]. The mechanisms that reduce NK activity in obesity are summarized in Fig. 4. Clinically, bariatric surgery improved NK cell activity in humans [239]. Glucagon-like-peptide-1 agonist (GLP-1) treatment in



### Fig. 4. Inhibition of NK cells in obesity.

In lean mice, mTOR signaling, P300, and MYC transcriptional proteins mediate the increase in cellular metabolism that is needed for NK polarization and the release of cytotoxic enzymes onto cancer cells. In obese mice, the uptake of lipids by NK cells leads to the formation of cytoplasmic lipid droplets (LDs) that inhibit mTOR signaling and destabilize P300 and MYC. This compromises cellular metabolism that is needed for the polarization and release of cytolytic enzymes by NK cells onto cancer cells. NK = Natural Killer, mTOR = mechanistic target of rapamycin, P300 = E1A Binding Protein P300, MYC = MYC Proto-Oncogene, BHLH Transcription Factor, GzmB = Granzyme B, PFN = Profilin 1, LD = lipid droplet.

humans also boosted NK activity [240]. These clinical findings are encouraging and suggest that moving to a low-fat diet and losing weight might be lifestyle changes that improve NK activity.

### 6.4. Myeloid-derived suppressor cells (MDSCs)

Chronic inflammation, infections, and cancer induce granulocytes and monocytes to develop into MDSCs [222,223]. MDSCs inhibit inflammation and suppress anti-tumor immunity [222,223]. Obesity-associated leptin expands the number of MDSCs in peripheral blood and the TME in mice [241]. Tumor-associated MDSCs express PD-L1, which inhibits the activity of tumor-associated CD8<sup>+</sup> T cells [241]. Clinically, obese women with TNBC had increased circulating MDSCs compared with lean women with TNBC [242]. In obese mice, renal tumors had increased MDSCs compared with renal tumors in lean mice [243]. Renal tumors in obese mice produced increased concentrations of C-C Motif Chemokine Ligand 2, which is an MDSC chemoattractant [243]. In obese mice, an increase in the number of MDSCs in the peritoneal cavity is associated with increased ovarian tumor burden [244]. In mice and humans, obesity-stimulated MDSC is associated with increased progression of oral squamous cell carcinoma [245]. In mice, obesity is broadly associated with an increase in the number of myeloid cells and a decrease in the number of lymphoid cells being produced in bone marrow [246]. In murine obesity, a subset of monocytes develops into APCs that activate neutrophils that travel to the lung and release neutrophil extracellular traps that "trap" metastatic mammary cancer cells in the lung [246]. This monocyte-neutrophil-based mechanism could, in part explain why breast cancer metastasis is increased in obesity [246].

### 6.5. Macrophages

Macrophages are the most abundant immune cell type in adipose

tissue and within the TME [247,248]. Macrophages respond to tissue signals to polarize towards an anti-inflammatory or proinflammatory activation state [249]. Macrophages in adipose tissue in obesity are classically M1 activated and release proinflammatory cytokines that drive chronic inflammation and cancer susceptibility [248]. Conversely, macrophages within the TME are alternatively polarized towards an M2 phenotype that produces cytokines that mediate antitumor immunity within the TME [247]. Consequently, TAMs that are M2-activated are associated with poor cancer prognosis [247]. Classically activated M1 macrophages also populate the TME and suppress cancer development by producing proinflammatory cytokines and presenting antigens to cytotoxic T cells [247]. Preclinical findings show a high-fat diet is associated with enhanced infiltration of M2-like macrophages and regulatory T cells into prostate tumors in mice [250]. Preclinical findings also show that high-fat diet-induced obesity correlates with upregulated expression of PD1 on TAMs, which reduces antigen presentation and phagocytic activity in murine MC38 colorectal tumors [251]. The obesity suppressive effects on TAMs were reversed by anti-PD1 antibody treatment [251]. Clinically, obesity is associated with an increase in the percentage of immunosuppressive M2-activated macrophages in breast adipose tissue [252]. These findings show that obesity regulation of TAMs promotes cancer development. The obesity signal that regulates TAMs, however, is unknown.

### 7. Cancer-associated adipocytes

Adipocytes in direct contact with cancer cells exhibit morphological, functional, and de-differentiation processes that are unique compared with adipocytes that are located far from the tumor—and because these changes are unique, adipocytes that are in direct contact with cancer cells are referred to as being cancer-associated adipocytes (CAAs) [253–255]. CAAs undergo lipolysis in the TME [253,255]. In CAAs, this is mediated by the activation of hormone-sensitive lipase by

cancer-secreted factors [254,256]. The induction of hormone-sensitive lipase in CCAs is partially mediated by adrenergic receptors in ovarian, but not breast cancer [254,256]. Free fatty acids are steadily liberated from lipid droplets by adipose tissue lipase expressed by cancer cells [254–257]. In cancer cells, free fatty acids are oxidized, which is not coupled to ATP synthesis but is essential for cancer cell invasiveness [256]. The flow of free fatty acids from adipocytes to cancer is enhanced in obesity [257].

In mice, in vivo adipocyte tracing studies show CAAs do not undergo apoptosis; instead, they transition to a fibroblast-like morphology [255]. This transition of an adipocyte to fibroblast-like cells is viewed as a de-differentiation process [253,255]. CAAs can re-differentiate to an adipocyte if removed from the tumor and placed in cell culture [255]. Thus, cancer-secreted factors maintain CAAs [255]. An interesting question is whether delipidation occurs first, which triggers de-differentiation, or if de-differentiation triggers delipidation, or the two processes are independent responses to direct interaction with cancer cells. In mice, genetically blocking adipocyte lipolysis prevents cancer-induced adipocyte de-differentiation in the context of mammary cancer [255]. This finding suggests that adipocyte delipidation leads to adipocyte de-differentiation in response to direct contact with cancer cells [255]. Mice with adipocytes that are refractory to lipolysis are also protected from tumorigenesis because implanted mammary tumors in these mice fail to grow and are significantly smaller than tumors implanted in mice with adipocytes that undergo lipolysis in response to being in direct contact with cancer cells [255]. Independent of lipid transfer, the de-differentiation of adipocytes in response to cancer cells might benefit the tumor by increasing total cellular heterogeneity within the TME [255] (Fig. 5). Specifically, cells that were adipocytes but are no longer adipocytes because they are in contact with cancer cells are a heterogeneous population of six different cell types in various stages of cellular transition that resemble inflammatory cells, macrophages, myofibroblasts, adipocyte progenitor cells, and mesenchymal cells in the murine TME [255] (Fig. 5). Single-cell RNA-seq studies show that these tumor-stimulated adipocyte-derived cells respond to hypoxia and alter pathways involved in the extracellular matrix and inflammation [255]. This heterogenous adipocyte response to cancer may be one mechanism by which de-differentiation of adipocytes promotes tumorigenesis by increasing cell heterogeneity in the tumor microenvironment [255] (Fig. 5). Targeting adipocytes in direct contact with cancer cells could foster new cancer therapies in obesity.

### 8. Conclusions and future directions

In obesity, chronic overfilling of adipocytes with lipids exceeds capillary support to adipose tissue, which in turn leads to the death of adipocytes that, in the process of dying, release chemokines and cytokines that recruit M1-stage macrophages to adipose tissue that mediate chronic low-level inflammation and skew T cell differentiation towards proinflammatory T<sub>H</sub>1 and T<sub>H</sub>17 T cells (Section 2). The inflammation is more severe in visceral adipose tissue. However, there is also an increase in low-grade inflammation in peripheral tissues that are prone to cancer, such as the breast, in obesity. There are many examples where obesity inhibits anti-tumor immunity, including exhaustion and lipid deprivation of CD8<sup>+</sup> T cells, inactivation of NK cells, and increased numbers of MDSCs (Section 6). Although obesity affects anti-tumor immunity through several mechanisms, it is not clear which perturbations are the most important for cancer progression and which mechanisms are cancer-type-specific. Many anti-tumor immunity mechanisms have been established in mouse models of obesity with correlates to human data. It will be essential to translate these findings to human cancer in obesity. The risk of cancer in obesity for many cancers is associated with a low hazard ratio [21]. Thus, obesity alone is not a promising biomarker for cancer. However, combining obesity with other cancer biomarkers might identify a subset of obese patients who are at high risk for aggressive cancer.





Cancer-associated adipocytes are in direct contact with cancer cells. Cancer cell-secreted factors act on adipocytes to induce lipolysis and the liberated FFAs are transferred to cancer cells for metabolism. Delipidation of adipocytes triggers de-differentiation into multiple cell types, including myofibroblasts, immune-like cells, and adipocyte progenitor cells that collectively increase cellular heterogeneity within the tumor microenvironment (TME), which promotes tumorigenesis. FFAs = Free fatty acids, TME = tumor microenvironment.

Given that obesity is a heterogeneous condition, a better understanding of which associated comorbidities drive T-cell dysfunction could lead to better-targeted interventions for cancer therapy. Identifying the obesity signal that drives T cell dysfunction could foster the development of novel therapeutic approaches to restore tumorassociated T cell function in obesity. Although the link between dysfunctional tumor-infiltrating T cells and obesity is relatively well characterized, less is known about how obesity regulates immunosurveillance of early cancer development. A more precise understanding of the mechanisms of T-cell responsiveness to immune checkpoint therapy in obesity may identify biomarkers that better predict patient response to immune-based treatment for cancer therapy [258]. Considering the sensitivity of tumor-associated T cells to diet, a better understanding of the mechanism could lead to therapies that incorporate diet to improve anti-tumor immunity [259].

### CRediT authorship contribution statement

**Cora E. Miracle:** Writing – review & editing, Writing – original draft. **Chelsea L. McCallister:** Writing – review & editing, Writing – original draft. **Richard D. Egleton:** Writing – review & editing, Writing – original draft. **Travis B. Salisbury:** Writing – review & editing, Writing – original draft.

### Declaration of competing interest

The authors declare they have no competing interests.

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