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Multifaceted Leptin network: the molecular connection between obesity and breast cancer

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Abstract

High plasma levels of leptin, a major adipocytokine produced by adipocytes, are correlated with increased fat mass in obese state. Leptin is emerging as a key candidate molecule linking obesity with breast cancer. Acting via endocrine, paracrine, and autocrine manner, leptin impacts various stages of breast tumorigenesis from initiation and primary tumor growth to metastatic progression. Leptin also modulates the tumor microenvironment mainly through supporting migration of endothelial cells, neo-angiogenesis and sustaining recruitment of macrophage and monocytes. Various studies have shown that hyperactive leptin-signaling network leads to concurrent activation of multiple oncogenic pathways resulting in enhanced proliferation, decreased apoptosis, acquisition of mesenchymal phenotype, potentiated migration and enhanced invasion potential of tumor cells. Furthermore, the capability of leptin to interact with other molecular effectors of obese state including, estrogen, IGF-1, insulin, VEGF and inflammatory cytokines further increases its impact on breast tumor progression in obese state. This article presents an overview of the studies investigating the involvement of leptin in breast cancer.

Introduction

Many prospective epidemiological studies have demonstrated that obesity, defined as excessive fat accumulation, is a pandemic condition that greatly influences risk, prognosis and progression of various cancers including breast cancer [1–7]. The Million Women study examined breast cancer incidence and mortality in relation to body mass index and reported that approximately half of the cancers can be attributed to obesity in postmenopausal women [8]. A study focused on invasive breast cancer reported that advanced grade and stage including lymph node metastases were more prevalent in obese women [9]. Investigating the relationship of obesity with mortality from breast cancer, many studies report that obese women in the highest quintile of body mass index (BMI) have double the death rate from breast cancer when compared with women in the lowest quintile [1,10–13]. In addition, in women with BMI in the highest quintile, an increased proportion of tumors were ER

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negative, had a high S-phase fraction, histological grade, mitotic cell count, expression levels of proliferation markers, and a larger tumor size [14]. Similar findings were reported by Daling et al., in a population-based follow-up study showing that the women in the highest quintile of BMI are more likely to have high histological grade, increased size and ER negative breast tumors [15]. Taking a molecular approach to examine the link between obesity and breast cancer, Creighton et al., examined the effect of patient obesity on gene expression of primary breast tumors. Transcription profiles were conducted on 103 tumors for which the patient's body mass index (BMI) was known and stratified into three groups according to BMI (normal, overweight and obese). Tumors from obese patients were compared to tumors from either normal or overweight patients, and the transcriptional level of 662 genes were found to be associated with the obese state defined as obesity-associated cancer gene signature. Obesity-associated cancer gene signature associated with worst patient outcome in meta-analysis [16]. Together, these studies have shown that obese women regardless of their menopausal status (in most cases) are likely to have metastatic breast cancer when they are first diagnosed, and to have a poor final outcome.

Several hypotheses have been proposed to explain this association and very popularly, particular emphasis has been placed on the increased production of estrogen from peripheral aromatization of androgens in adipose tissue. High levels of estrogen can promote the development of post-menopausal carcinoma of the breast but despite the relationship between obesity and high estrogenic activity, it has become evident that this cannot fully explain the associations between body weight and breast cancer risk and poor prognosis because, in studies of both premenopausal and postmenopausal women, large subgroups of obese women were identified with ER-negative breast cancer exhibiting rapid growth and aggressive metastatic biological character. According to the newer hypothesis, obesity is now considered as an endocrine disorder and is associated with increased circulating levels of insulin, bioavailable insulin-like growth factor (IGF-1), inflammatory cytokines, vascular integrity factors such as vascular endothelial growth factors (VEGF), plasminogen activator inhibitor (PAI)-1 and adipocytokines [10,11,17]. These mediators, their interacting partners and pathways form the complex molecular network by which obese state impacts pathological manifestation of carcinogenesis.

A typical feature of obese state is an excess fat mass as a consequence of hypertrophy and hyperplasia of adipocytes. With recent research elevating the status of adipocytes from 'inert energy storing cells' to 'active endocrine organs', there has been significant interest in the potential role of adipocytokines (a group of proteins synthesized in adipose tissues) in the development of obesity-associated cancers. Adipocytokines are mainly produced by the adipocytes and the stromal cells (fibroblasts) that can potentially differentiate to mature adipocytes and macrophages that infiltrate the adipose tissue. Acting by endocrine, paracrine, and autocrine mechanisms, adipocytokines affect various biological processes. [18,19]. Among various adipocytokines, leptin, owing to its myriad oncogenic effects on carcinogenesis have come to be recognized as an important mediator of molecular effects of obesity. In this review, we will focus on adipocytokine leptin, its normal role in breast, clinical and preclinical studies showing its association with breast cancer, notably its effect on breast cancer cells growth and metastatic properties. We will also emphasize the interaction of leptin with other important mediators of obesity-breast cancer axis with the

hypothesis that only by studying the interconnected network of various biological mediators of obesity can a complete understanding be achieved.

Dysregulated Leptin Levels in Obesity and Breast Cancer

Leptin, a product of the obese (*ob*) gene, is a neuroendocrine hormone that has attracted attention since its identification in 1994 [20] as it launched a new field in obesity research. Leptin is synthesized and secreted in proportion to body mass predominantly from preadipocytes and adipocytes, and to a lesser extent from placenta, stomach and skeletal muscle. Circulating as a 16 kDa non-glycosylated protein partially bound to plasma proteins, leptin targets hypothalamus, and peripheral organs, including liver, skeletal muscle and pancreas [21,22]. In addition to its role as a satiety hormone research over the last few years provided important clues about its apheliotropic actions, its role in the pathogenesis of atherosclerotic vascular disease and importantly carcinogenesis [23–25]. Biological actions of leptin are mediated through binding to the extracellular domain of specific membrane receptor (leptin receptor) present in a variety of tissues localized to the cell membranes [26]. Leptin receptors are characterized as class I cytokine receptors typically containing a cytokine receptor homologous domain in the extracellular region [27]. Up to now, six isoforms of the leptin receptor have been identified. All isoforms have a similar extracellular ligand binding domain at the amino terminus but they differ at the intracellular carboxy-terminal domain. While all five short isoforms have transmembrane domains, only the long form has the intracellular motifs necessary for activation of signaling pathways [28].

Although leptin was found as an afferent satiety signal, regulating appetite and energy expenditure in both humans and rodents [29], obese state is not related to deficiency of leptin. In fact, high plasma levels of leptin are correlated with increased fat mass in obese state and decreased levels are found in lean humans and animals [30,31]. Multiple epidemiological studies have examined serum leptin levels in women with breast cancer over past few years. The relationship of circulating leptin levels with breast cancer appears to be complex with most studies reporting a clear positive association between increased serum leptin levels and elevated breast cancer risk while few others found no change or even reduced levels of leptin associated with breast cancer [32–35]. The contradiction in these results can be largely attributed to confounding factors such as disease stage, inclusion of small number of subjects, considering only premenopausal subjects or combining pre and post menopausal women. To address the contradictory results that have been reported regarding the association between leptin level and breast cancer, Niu et al., recently performed a meta-analysis utilizing data from 23 relevant studies involving 2058 breast cancer patients, 2078 healthy controls and 285 breast benign controls. They concluded that the circulating leptin levels were lowest in healthy people while increasing levels were found in different groups as follows: healthy people < breast benign diseases patients < breast cancer patients < lymph node metastasis positive patients [36]. Clear evidence has emerged for postmenopausal breast cancer patients with ER positive breast cancer where serum leptin levels significantly correlated with poor clinicopathological tumor classification [37]. A case control study investigating the relationships of plasma leptin level and anthropometric measures of adiposity with the risk of breast cancer reported that overall higher leptin concentrations were significantly associated with an increased risk of breast cancer. They

also found that the associations of leptin with breast cancer risk persisted after adjustment for obesity indices suggesting that leptin may have an independent role in breast tumorigenesis [38]. Another study analyzing leptin and leptin receptor involvement in breast carcinoma showed positive correlation with tumor size [39]. Leptin signaling and breast cancer were linked in a clinical study showing that leptin receptors were not detectable in normal mammary epithelial cells by immunohistochemistry, whereas in 83% of cases, carcinoma cells showed positive staining for the leptin receptor [40]. Of importance, overexpression of leptin was observed in 92% of breast tumors examined but in none of the cases of normal breast epithelium. The positive correlation of both leptin and leptin receptor suggested that leptin can potentially act on breast cancer cells via an autocrine pathway [40]. Another important study reported a positive relationship between blood leptin levels and breast cancer risk. Also, the degree of leptin mRNA expression in the peritumoral adipose tissue was found to be significantly higher in the breast cancer patients than the control women [41]. Studies also indicated that leptin and leptin receptor are overexpressed in primary and metastatic invasive ductal breast carcinoma compared with non-cancer mammary tissue [42]. In a recent study, 78 obese patients with newly diagnosed breast cancer and without diabetes and 78 obese women without breast cancer and diabetes were evaluated for their leptin levels. Higher leptin levels were found in women with breast cancer and obesity in comparison to similar obese women without breast cancer (Romero-Figueroa, et al., *Clinical Breast Cancer*, 2013, *In press*).

Evidence for the Role of Leptin in Mediating Obesity-Breast Cancer Link

In breast microenvironment, adipokine leptin is secreted by adipocytes which form the bulk of the human breast being the most abundant cell type surrounding breast cancer cells and mammary epithelial cells which also produce leptin [43]. Several preclinical animal studies have also put forth the important role of leptin in breast tumorigenesis. *In vivo* studies examining the impact of leptin-axis on breast carcinogenesis utilizing genetic loss-of-function mutants for leptin or the leptin receptor showed that leptin or leptin receptor-deficient MMTV-Transforming growth factor- α (MMTV-TGF- α /Ob/Ob and MMTV-TGF- α /db/db) mice did not develop oncogene-induced mammary tumors [44,45]. These studies suggested that intact leptin-axis is required for spontaneous mammary tumorigenesis. In another approach, when hypothalamic Ob-Rb (long-form leptin receptor) reconstituted db/db (LEPR-null) mice ($db/db^{Nse+/+}$) [46] were crossed with MMTV-PyMT mice, it was found that an Ob-Rb-mediated signaling promoted mammary tumor growth and metastasis [47]. Physiologically relevant high-fat diet-induced obese mouse models have also been used for manipulating leptin levels *in vivo*. Obese MMTV-TGF α mice certainly showed higher levels of leptin as well as accelerated growth of mammary tumors [48]. In contrast, obesity and elevated leptin levels did not affect the development of estrogen-negative breast tumors in MMTV-neu mice [49]. When xenografts of MMTV-Wnt1 tumors were transplanted in diet-induced obese mice, the tumors developed faster as compared to lean counterparts [50]. In coherence with the important role of leptin in breast tumorigenesis, xenografts of MMTV-Wnt1 cancer cells transplanted into leptin-deficient obese (*Ob/Ob*) mice displayed stunted growth [51]. In athymic-nude mice xenograft studies, leptin treatment significantly increased breast tumor growth [52]. Genetically obese Zucker rats

which have a leptin receptor defect and lean, non-litter mates were injected with a chemical carcinogen methylnitrosourea and analyzed for the development of breast carcinoma. A smaller percentage of obese Zucker rats developed carcinomas as compared to the lean, non-litter mates [53]. High level of leptin circulating in the blood in obese state exerts its biological effects on responsive cells in a classical endocrine manner. In addition, produced by the adipocytes in the breast tumor microenvironment, leptin acts on breast cancer cells through paracrine pathways. It is important to note that breast cancer cells possess leptin receptors and secrete leptin which acts in an autocrine manner (Figure 1). The studies discussed above suggest the importance of endocrine, paracrine and autocrine effects of leptin and the importance of leptin receptor in breast carcinogenesis.

Interaction of Leptin with Other Important Mediators of Breast Cancer

Leptin has been shown to interact with various important mediators of breast cancer in obese state including estrogen signaling network, insulin, IGF-1, angiogenic pathways and inflammatory cytokines impacting breast cancer initiation, growth and metastatic progression (Figure 2).

Estrogen and Aromatase: functional crosstalk with leptin

Estrogen plays an integral role in normal mammary gland development and is associated with breast tumor progression [54]. In postmenopausal women, adipose tissue serves as a major source of estrogens primarily produced by aromatization of androstenedione via cytochrome P450 19A1 (CYP19A1) also known as aromatase [55]. Functional crosstalk between leptin and estrogen signaling network further contributes to breast carcinogenesis exemplifying the interactions between resident adipocytes and breast epithelial cells. Leptin enhances aromatase mRNA expression, aromatase content, and its enzymatic activity in breast cancer cells in an ERK and Stat3-dependent manner as the presence of MAPK inhibitor, ERK2 dominant negative or Stat3 dominant negative mitigates the stimulatory effects of leptin [56]. Leptin-induced aromatase activity further leads to increased production of estradiol and estrogen receptor (ER) signaling in breast cancer cells [56]. Another mechanism by which leptin increases estrogen signaling in breast cancer cells is via direct transactivation of ER in the absence of cognate ligand. Leptin treatment induces classical features of ER activation including nuclear translocation, downregulation of ER mRNA and protein and upregulation of classical ER responsive genes which is abrogated in the presence of ERK inhibitor or dominant negative ERK2 [57]. It has been shown in multiple cancer cell lines that leptin activates phosphorylation of ERK [58–60]. ER is known to get phosphorylated on Ser-118 by ERK in response to upstream signaling events including estrogen, insulin/insulin like factor1 (IGF-1) and this phosphorylation is required for functional activation of ER [61]. In a feed-forward mechanism, leptin activates ERK which in turn phosphorylates ER resulting in its activation in breast cancer cells. Leptin mediated activation of ERK and Stat3 has also been associated with increased expression levels of ER in MCF7 breast cancer cells [62]. Leptin-mediated increased expression of ER was lost upon silencing of leptin receptor indicating a functional dependence. Analysis of breast cancer samples from 33 patients at different stages of disease showed a statistically significant correlation between the expression of leptin receptor and estrogen receptor [62]. Exogenous leptin treatment has been reported to increase ER expression in breast tumors in

nude-mouse xenograft model [63]. Chronic treatment of breast cancer cells with leptin potentiates the mitogenic actions of estrogen and alters the ratio of ER α to ER β [64] indicating that high levels of leptin in breast tumor microenvironment can enhance estrogen-signaling. According to these findings, leptin plays an important role in modulating E2-ER network in breast cancer cells. On one hand, leptin enhances estrogen levels by increasing the conversion of androstenedione to estrogen especially in post-menopausal conditions and on the other hand activates ER function in a ligand-independent manner. Interestingly, estrogen-signaling also in turn modulates leptin function as overexpression of estrogen receptor increases leptin-induced Stat3 activity in breast cancer cells [65]. Together, these studies indicate a biologically important crosstalk between estrogen and leptin signaling networks.

Connection between insulin, IGF1 and leptin

Hyperinsulinemia or insulin resistance is associated with obese conditions and increases the risk and progression of breast cancer [66]. Insulin, a peptide hormone produced by the beta cells of the pancreas and released in response to increased blood glucose, is reported to exert antiapoptotic effects at normal levels while supraphysiological levels of insulin, as evident in obese state, have mitogenic effects. Insulin mediates its tumor-promoting effects directly via the insulin receptor (IR) or insulin-like growth factor 1 receptor (IGF-1R). Elevated insulin levels induce extracellular-signal regulated kinase (ERK) and phosphatidylinositol-3 kinase (PI3K) pathways in breast cancer cells leading to downstream activation of several pathways and oncogenic processes [67]. Studies examining the relationship between obesity related stimuli including insulin and leptin in breast cancer reported a positive correlation between leptin and insulin as high concentrations of insulin stimulated leptin mRNA in both ER α -positive MCF-7 and ER α -negative MDA-MB 231 cell lines [42,68]. Elevated level of IGF-1, a peptide growth factor that shares ~50% sequence homology with insulin, observed in obese state is positively associated with increased breast cancer risk [69]. IGF-I receptors are known to mediate the proliferative effects of insulin as the insulin receptor (IR) and the receptor for IGF-I (IGF-IR) exhibit more than 50% of overall sequence homology and the tyrosine kinase domain of the α -subunit share 84% homology. Also, their respective ligands, insulin, and IGF-I share 40–50% homology supporting interaction of insulin and IGF-1 with both IR and IGF-1R [66]. IGF-1 overexpression leads to excessive proliferation and survival signals for the breast tumor development [70]. IGF-1R is overexpressed in ~50% of primary breast tumors compared with normal tissue indicating that these carcinomas have enhanced responses to the mitogenic and anti-apoptotic effects of IGF-1 [71] and inactivation of IGF-1R results in reduced breast tumor growth and metastasis *in vivo* [72]. Studies from our group and others have shown interactions between leptin signaling network and IGF-1 [73,74]. We demonstrated that combined treatment with leptin and IGF-1 significantly increased proliferation as well as invasion and migration of breast cancer cells. A novel bidirectional crosstalk between leptin and IGF-1 signaling was found; IGF-1 treatment induced remarkable phosphorylation of leptin receptor (Ob-Rb) and leptin treatment induced tyrosine phosphorylation of IGF-1 receptor (IGF-1R) while co-treatment induced synergistic phosphorylation of both receptors and association of Ob-Rb and IGF-1R along with activation of downstream effectors, Akt, ERK, IRS-1, and IRS-2. Interestingly, the biological effects of this crosstalk were mediated by epidermal growth factor receptor

(EGFR) activation depending on proteolytic release of EGFR ligands as broad-spectrum matrix metalloproteinase inhibitor, GM6001 could inhibit this effect. Our study further showed that inhibition of EGFR activation using clinically relevant EGFR inhibitors, erlotinib and lapatinib, inhibited leptin and IGF-1 induced invasion and migration of breast cancer cells [74]. Together, these studies suggest that interactions between leptin, insulin and IGF-1 could contribute to breast cancer initiation, growth and metastatic progression in obese state.

Impact of leptin on angiogenic network

Vascular endothelial growth factor (VEGF), possessing angiogenic, mitogenic, and vascular permeability-enhancing activities specific for endothelial cells, is a key angiogenic factor [75]. VEGF is considered an adipokine due to its synthesis in adipose tissue and is positively correlated with increased BMI [76]. High circulating levels and increased tumoral expression of VEGF is associated with poor prognosis [77,78]. Recent studies have shown that leptin is a pro-angiogenic cytokine promoting angiogenesis, upregulating VEGF and VEGF receptors in breast cancer. Leptin induces angiogenesis in normal rat corneas but not in corneas of rats genetically lacking leptin receptors [79] indicating the involvement of leptin-signaling network. Using various *in vitro* and *in vivo* approaches, various studies have shown that leptin increases proliferation of human umbilical venous endothelial cells with a potency comparable to VEGF treatment, induces capillary-like tube formation and stimulates angiogenesis in chorioallantoic membrane (CAM) assay and disc implantation assay [80–82]. Synergistic effects of leptin, VEGF and bFGF have also been shown to stimulate angiogenesis [83]. Leptin directly upregulates VEGF in breast cancer cells via mediating various canonical and non-canonical signaling pathways [84,85]. Leptin-mediated increased Stat3, ERK, and Akt phosphorylation leads to upregulation of VEGF mRNA and protein in mouse mammary cancer cells (MT) and inhibition of these canonical signaling pathways using specific small-molecule inhibitors abrogates the effect of leptin on VEGF levels. Leptin-mediated VEGF upregulation also involves modulation of HIF1 α and NF- κ B and inhibitors of these transcription factors such as NS398 (for HIF1 α) and inhibitor IKK antagonist (for NF- κ B) negatively impact leptin-mediated VEGF expression [84]. Leptin treatment has also been reported to increase the expression of VEGF receptor 2 (VEGFR2) independent of VEGF [86] resulting in promotion of mammary tumor growth. Using multiple breast cancer cells, Ray et al., show that leptin treatment results in elevated levels of VEGF [87]. Upregulation of VEGF and VEGFR2 by leptin have important implications in breast carcinogenesis as neoangiogenesis is required to fulfill the nutrients and oxygen needs of growing tumors. In addition to modulating VEGF/VEGFR2 axis to support neoangiogenesis, leptin induces the expression of several other molecules including MMP-2, MMP-9, leukemia inhibitory factor (LIF), interleukin-1 (IL-1), and also α v β 3 integrin in various model systems to further increase tumor angiogenesis. Leptin stimulates MMP-2 activity in breast cancer cells via Jun N-terminal kinase activation [88]. Leptin also regulates IL-1, LIF and their respective receptors, IL-1R tI and LIFR in benign (primary and HES) and cancerous-endometrial epithelial cells (An3Ca, SK-UT2 and Ishikawa), and induces a greater increase in LIF levels in cancer as compared to benign cells [89]. Furthermore, leptin modulates fibroblasts and macrophages to stimulate increased secretion of pro-angiogenic factors. In tumor microenvironment, breast tumors produce a high level of IL-1 *in vivo*

which triggers leptin expression in stromal cells and infiltrating immune cells [90]. Leptin is also shown to induce several signaling pathways for transcriptional and translational expression of IL-1 and associated components in breast cancer cells and inhibition of IL-1 abrogates leptin-induced VEGF expression [91]. Further research is needed to delineate the molecular mechanisms by which leptin crosstalks with various components of angiogenic network and exerts its effect on promotion of angiogenesis in breast cancer.

Leptin and inflammatory cytokines

Inflammatory cytokines such as tumor necrosis factor- α (TNF- α) and IL-6 are primarily produced by infiltrating macrophages while small amounts are synthesized by adipocytes in obese state. Increased plasma concentration of TNF- α , IL-17 and IL-6 are associated with high BMI levels [92,93]. Tumor associated macrophages express multiple receptors including leptin-receptor (Ob-R), toll-like receptor (TLR4), and play an important role in breast tumor progression mediating crosstalk between tumor cells and adipocytes [94]. Adipocytes secrete adipokine leptin which binds to its receptor on macrophages and stimulate the secretion of multiple pro-inflammatory and pro-angiogenic cytokines such as IL-1, TNF- α , IL-6, IL-11 and nitric oxide modulating macrophage phenotype (anti-inflammatory M2 and pro-inflammatory M1). M1-polarized or classically activated macrophages colonize adipose tissue [95]. Adipose tissue macrophages exhibit increased inflammatory properties during diet-induced obesity [96]. Several studies have reported leptin-mediated modulations of proinflammatory cytokines in multiple systems. Leptin treatment induces IL-17 expression in CD4⁺ cells while macrophages are shown to produce IL-6 and TNF- α in response to leptin. Dendritic cells exhibit increased production of multiple cytokines including IL-1, IL-6, IL-12, TNF- α and MIP-1 α upon leptin stimulation [97–99]. Leptin upregulates IL-1 and TNF- α in macrophages and tissue factor (TF) in breast cancer cells via TNF- α [100,101]. Although high circulating levels of some cytokines and leptin are known to be negative prognostic markers for breast cancer and some studies have shown interactions between them, further studies are needed to clarify their cell-specific molecular link and importance in breast tumor progression.

How Does Leptin Influence Breast Cancer Growth and Metastasis?

Activation of oncogenic pathways: multipartite leptin network

Various research groups have been studying the oncogenic role of leptin in breast cancer. *In vitro* studies from our lab and others have shown that leptin regulates various molecules and oncogenic pathways involved in proliferation, adhesion, invasion, migration, inflammation and angiogenesis, such as cyclin D1, survivin, β 3 integrin, interleukin-1 (IL-1), IL-1 receptor, vascular endothelial growth factor and its receptor type 2 in breast carcinogenesis [52,58,74,86,102–106]. Concomitant activation of multiple signaling pathways including ERK, Akt and Stat3 network via leptin has been reported in breast cancer [58,59,74,102,107]. Leptin modulates the expression of various important genes implicated in breast carcinogenesis via Stat3 activation. It is interesting to note that leptin can directly manipulate the transactivation function of certain coactivator molecules leading to alterations in local chromatin structure. For example, leptin induces cyclin D1 expression increasing histone acetylation at the cyclin D1 promoter. Genes in repressed state have been

shown to be associated with methylation of K9-dimethylated H3 whereas the active state is associated with increased methylation of K4-dimethylated H3 [108]. Leptin increases H3-K4 methylation and decreases H3-K9 methylation creating a permissive environment for the recruitment of specific coactivator complexes including Med1 and SRC1 resulting in increased breast cancer cell growth [102]. Leptin also regulates cyclin D1 expression in several other breast cancer cells [87,109,110] and luminal epithelial cells of mouse MMTV-Wnt1 mammary tumors [111]. Leptin-mediated increased expression of survivin, a member of inhibitor-of-apoptosis proteins (IAP) family, has been reported in breast cancer cells [52,112,113]. Our recent work showing leptin-mediated increased survivin expression implicates a complex upstream network involving activation of EGFR-Notch1 axis. These studies show that leptin activates Notch1 and induces recruitment of NICD (transcriptionally active intracellular Notch) to survivin promoter. Interestingly, Notch 1 activation by leptin is mediated via epidermal growth factor receptor (EGFR) transactivation implicating another oncogenic pathway in leptin network [52]. In addition, involvement of EGFR transactivation in leptin function has also been reported in gastric cancer cells [114], and oesophageal adenocarcinoma cells [115]. Several studies have reported multiple crosstalks between leptin and other contributors of mammary tumorigenesis and progression including IGF1, IL6, Notch and sex hormones [52,62,74,116] resulting in enhanced growth and metastatic properties of breast cancer cells. Hence, highly-active leptin-induced signaling-network (Figure 3) could contribute to various aspects of breast tumorigenesis and metastasis by modulating various molecular mediators and key pathways.

Leptin, epithelial-mesenchymal transition and breast cancer stem cells

Epithelial mesenchymal transition (EMT) is a crucial step in the induction of cell motility and enhanced survival during physiological processes such as embryonic development and wound healing as well as in pathological situations like malignant cells undergoing invasion and metastasis [117]. Recent studies from our lab present a central role of leptin in acquisition of mesenchymal characteristics and aggressive behavior in breast cancer. Leptin treatment induces breast cancer cells to undergo a transition from epithelial to spindle-like mesenchymal morphology. The key mechanism to account for this important function of leptin is that it increases β -catenin stabilization and nuclear translocation. Leptin-mediated stabilization of β -catenin is achieved by concomitant activation of Akt/GSK3 and MTA1/Wnt1 signaling leading to destruction of LKB1-GSK3 β -Axin complex. We also provide molecular evidence supporting the regulatory role of MTA1 in leptin-induced Wnt1 upregulation, β -catenin stabilization and nuclear translocation. These studies implicate a previously unrecognized crosstalk between leptin and MTA1/Wnt signaling in epithelial-mesenchymal transition of breast cancer cells [118]. Two other studies report involvement of leptin-signaling in EMT: role of leptin in endocardial cushion formation by modulating EMT [119] and activating hedgehog pathway to play a role in EMT in hepatic stellate cells [120]. Given that EMT plays an integral role in metastatic progression of breast tumors as well as sustaining the breast cancer stem cells (BCSCs), it is of interest to discuss the current knowledge regarding the impact of leptin on breast cancer stem cells.

BCSCs, known for their characteristic ability to undergo self-renewal as well as tumor differentiation, play an important role in breast cancer initiation, growth and metastatic

manifestation. Several signaling pathways including Notch, Hedgehog, Wnt/ β -catenin and TGF- β play an essential role in stem cell function during embryogenesis as well as oncogenesis [121]. It is interesting to note that leptin has been reported to regulate many signaling pathways and transcription factors implicated in BCSCs (discussed in previous section). Recently, Zheng et al. observed that MMTV-Wnt1 xenografts grow poorly in leptin knockout ob/ob mice in comparison to wild-type mice. Analysis of these tumor samples led to the identification of a leptin-responsive cell population that was present only in the tumors from wild-type mice. Fluorescent-activated cell sorting (FACS) for CSC markers including CD29 (integrin β 1), CD49f (integrin α 6), and CD24 (heat stable antigen) show that CD29⁺CD24⁻ and leptin receptor expressing CSC population is highly increased in the presence of leptin. These studies indicate the involvement of leptin in CSC survival [51]. Utilizing leptin-deficient ob/ob and leptin-receptor-deficient db/db, another study show that leptin-receptor (Ob-R) is a characteristic feature of tumor-initiating stem cells (TISCs) and is regulated by the core pluripotency-associated transcription factors OCT4 and SOX2. Also, TISCs demonstrate increased phosphorylation of pluripotency-associated oncogene Stat3 and induction of OCT4 and SOX2 in response to leptin [122]. It is suggested that leptin receptor plays an important role in the expression of Nanog, OCT4 and SOX2 in BCSCs, viability of BCSCs, proliferation and tumor-initiating activity [123]. These findings raise the possibility that obesity and its associated increased level of leptin-network can act as a mechanistic link between obesity, increased maintenance of cancer stem cells, augmented cancer recurrence, increased distant metastasis and overall poor survival. Leptin signaling has also been implicated in mediating interactions between K303R mutant estrogen receptor-expression breast cancer cells and cancer-associated fibroblasts (CAFs). It is proposed that leptin plays an integral role in a bidirectional crosstalk between breast cancer cells and the “educated” CAFs driving tumor progression [124].

Concluding Remarks

Breast cancer progression, a multistep process involving tumor initiation, primary tumor growth, invasion, metastatic progression, involves complex interaction with various stromal components including endothelial cells, immune cells, fibroblasts and adipocytes. Adipocytes are the major component in breast cancer microenvironment and are known to secrete various adipokines. Adipokine leptin is produced by adipocytes as well as breast cancer cells and acts in a paracrine, autocrine as well as endocrine manner. Various studies over the past few years have shown that leptin impacts breast tumor progression directly by interacting with breast tumor cells and indirectly by influencing various components of the tumor microenvironment. Hyperactive leptin-signaling network during obese state leads to concurrent activation of multiple oncogenic pathways resulting in: enhanced proliferation, decreased apoptosis, acquisition of mesenchymal phenotype, potentiated migration and enhanced invasion potential of tumor cells. In addition, leptin supports neo-angiogenesis and sustained recruitment of macrophages and monocytes, which upon leptin-stimulation, secrete VEGF and proinflammatory cytokines. Leptin also crosstalks with several other molecular effectors of obese state including, estrogen, IGF-1, insulin, VEGF and inflammatory cytokines, many times, potentiating their activities. Recent research advances have also implicated leptin in cancer initiation and breast cancer stem cells though further research is needed to strengthen these conclusions and elucidate the underlying molecular

mechanisms. Given all the potential roles of leptin in various steps of breast cancer progression and its strong link with obesity, the leptin-signaling network emerges as an attractive therapeutic target for the obese breast cancer patients.

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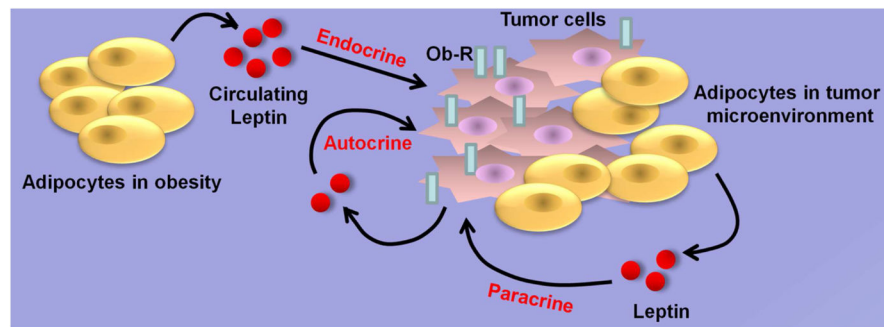


Figure 1. Endocrine, paracrine and autocrine actions of leptin

Leptin circulates at high levels in obese conditions and impacts breast tumor growth in an endocrine manner. Resident adipocytes secrete leptin in breast tumor microenvironment and act on the infiltrating breast cancer cells exhibiting paracrine effect of leptin. A direct autocrine effect of leptin is observed when leptin secreted from breast tumor cells impacts tumor growth acting via leptin receptors present on the breast tumor cells.

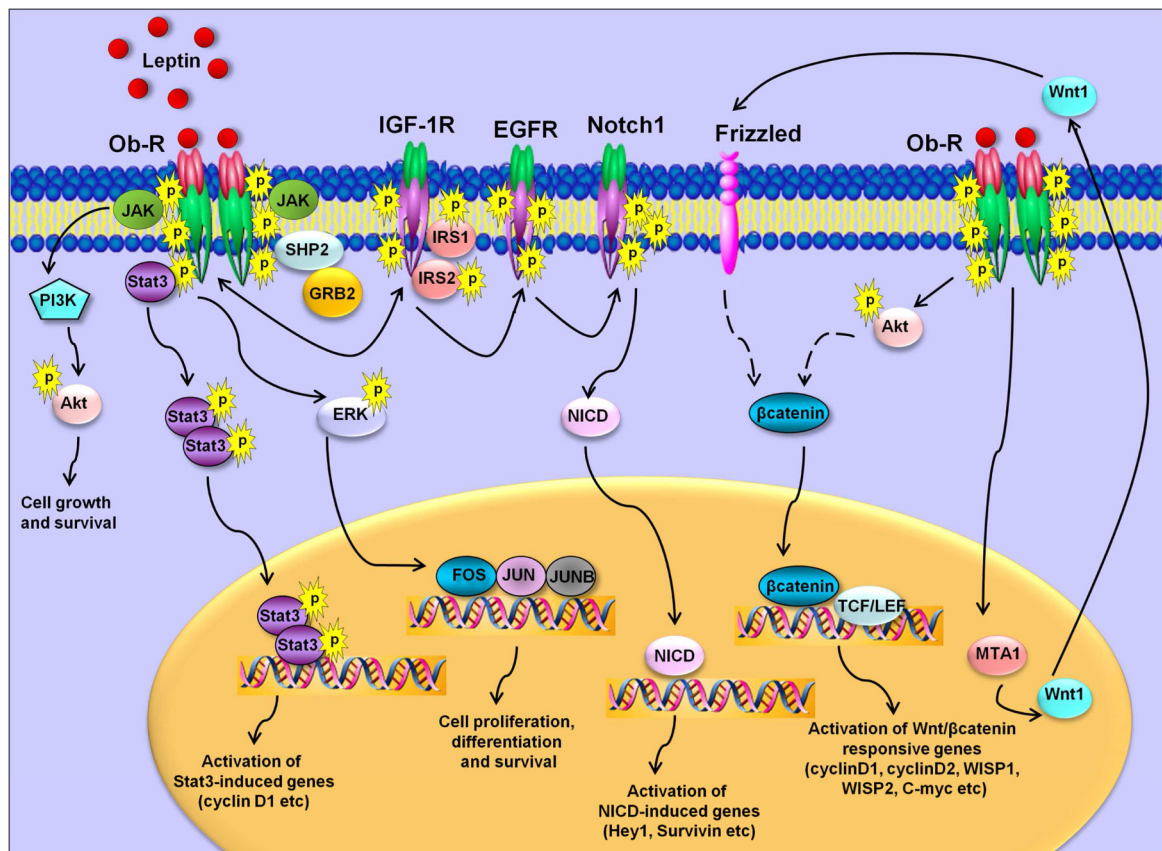


Figure 2. Leptin interacts with various important mediators of breast cancer

Leptin produced by adipose tissue binds to leptin receptor expressing cells in the tumor microenvironment including breast cancer cells, breast cancer stem cells (BCSCs), endothelial cells, fibroblasts and immune cells. Leptin activates multiple oncogenic pathways and impacts various steps of tumor progression.

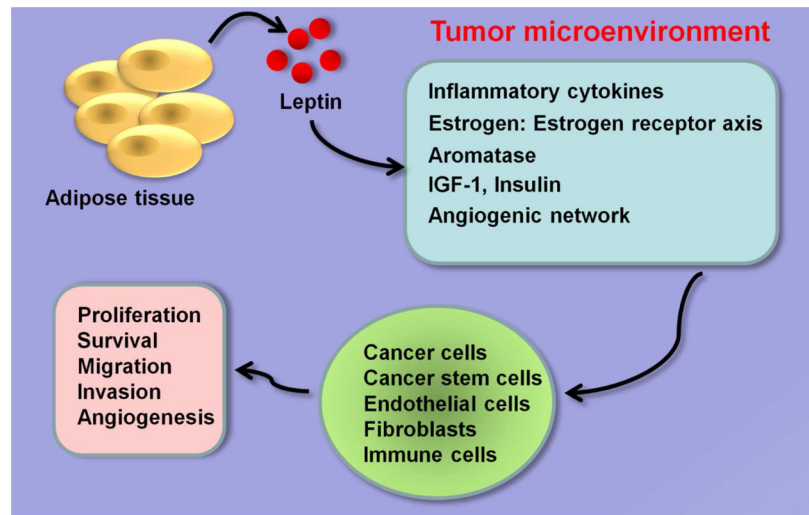


Figure 3. Complex leptin signaling network

Leptin binds to long-form of leptin receptor (Ob-R) resulting in conformational changes and receptor oligomerization. Leptin receptor gets phosphorylated followed by JAK activation which in turn phosphorylates Ob-R. These early events trigger activation of multiple pathways such as Akt activation, ERK phosphorylation and Stat3 activation. Bidirectional crosstalk occurs between Ob-R and IGF-1R. Leptin also transactivates EGFR and activates Notch1 resulting in release of NICD. Leptin increases the expression of MTA1 resulting in increased levels of Wnt1 and activation of Wnt1/ β -catenin network. Abbreviations: AKT, protein kinase B; GRB2, growth factor receptor-bound protein 2; JAK, Janus kinase; Ob-R, leptin receptor; MAPK, mitogen-activated protein kinase; PI3K, phosphatidylinositol 3 kinase; SHP2, Src homology 2-containing tyrosine phosphatase; STAT3, signal transducer and activator of transcription 3; insulin-like growth factor 1 receptor, IGF-1R; epidermal growth factor receptor, EGFR; transcriptionally active intracellular Notch, NICD; metastasis associated protein 1, MTA1.