Benzyl isothiocyanate inhibits oncogenic actions of leptin in human breast cancer cells by suppressing activation of signal transducer and activator of transcription 3

Su-Hyeong Kim¹, Arumugam Nagalingam², Neeraj K. Saxena²,³, Shivendra V. Singh¹ and Dipali Sharma²,⁴,*

¹Department of Pharmacology and Chemical Biology, and University of Pittsburgh Cancer Institute, University of Pittsburgh School of Medicine, Pittsburgh, PA 15213, USA, and ²Department of Hematology and Medical Oncology, and ³Department of Medicine, and ⁴Department of Medicine, and Emory Winship Cancer Institute, Emory University School of Medicine, Atlanta GA 30322, USA

To whom correspondence should be addressed. Dipali Sharma, Department of Oncology, Sidney Kimmel Comprehensive Cancer Center at Johns Hopkins, 1650 Orleans Street, CRB 1, Room 145, Baltimore, MD 21231-1000, USA. Tel: +1 410 455 1345; Fax: +1 410 614 4073; Email: dsharma7@jhmi.edu

Correspondence may also be addressed to Shivendra V. Singh: Tel: +1 412 623 3263; Fax: +1 412 623 7826; Email: singhs@upmc.edu

Molecular effects of obesity, a well-established risk factor for breast cancer progression, are mediated by adipocytokine leptin. Given the important role of leptin in breast cancer growth and metastasis, novel strategies to antagonize biological effects of this adipocytokine are much desired. We showed previously that benzyl isothiocyanate (BITC), a constituent of edible cruciferous vegetables (e.g. garden cress), confers significant protection against mammary carcinogenesis in a transgenic mouse model. The present study provides first evidence for the efficacy of BITC against oncogenic effects of leptin. The BITC treatment circumvented leptin-induced clonogenicity and anchorage-independent growth of MDA-MB-231 and MCF-7 human breast cancer cells. Leptin-stimulated migration and invasion of these cells was also inhibited in the presence of BITC. Analysis of the underlying molecular mechanisms revealed that BITC treatment suppressed leptin-induced Stat3 phosphorylation and cyclin D1 transactivation. The BITC-mediated inhibition of MDA-MB-231 xenograft growth correlated with a modest yet significant decrease in levels of Tyr705 phosphorylated Stat3. The BITC treatment efficiently inhibited Stat3 and SRC1 recruitment to cyclin D1 promoter in a chromatin immunoprecipitation analysis. Furthermore, overexpression of constitutively active Stat3 imparted significant protection against BITC-mediated inhibition of cyclin D1 transactivation, whereas RNA interference of Stat3 resulted in a significant increase in BITC-mediated inhibition of cyclin D1 transactivation in the presence of leptin. These results indicate that Stat3 plays an important role in BITC-mediated inhibition of leptin-induced cyclin D1 transactivation. In conclusion, BITC could potentially be a rational therapeutic strategy for breast carcinoma in obese patients with high leptin levels.

Introduction

Obesity is fast becoming a pandemic in the USA. Studies examining the relationship of obesity with mortality from breast cancer have found that obese women in the highest quintile of body mass index have double the death rate from breast cancer when compared with women in the lowest quintile (1,2). In addition, in women with body mass index in the highest quintile, the increased proportion of tumors were estrogen receptor (ER) negative, had a high S-phase fraction, histological grade, mitotic cell count, expression levels of proliferation markers and a larger tumor size.

These clinical observations cannot be explained only by higher estrogen levels that are associated with obesity. Importantly, independent of their menopausal status, obese breast cancer patients exhibit a higher risk for lymph node metastasis, larger tumor burden and higher mortality when compared with non-obese breast cancer patients. Obesity is associated with an increase in number and size of adipocytes that greatly alters the local and systemic secretion of biologically active polypeptides, adipocytokines such as leptin (3). Adipocytokines affect various biological processes acting by endocrine, paracrine and autocrine mechanisms (3).

Leptin was found as an afferent satiety signal, regulating appetite and energy expenditure in both humans and rodents (4). Leptin was recognized only as a satiety hormone for many years until research over the last few years provided important clues about its anabolicotropic actions, its role in the pathogenesis of atherosclerotic vascular disease (5,6) and importantly carcinogenesis (7). Leptin circulates as a 16 kD protein partially bound to plasma proteins and exerts its actions through specific cell surface receptors [leptin receptor (LR)] present in a variety of tissues localized to the cell membranes (8). Several epidemiological studies have linked high levels of plasma leptin with increased risk for breast carcinogenesis, but a direct link between leptin signaling and breast cancer was first established in a clinical study by Nagawa group in 2004 (9). They found that LRs were not detectable in normal mammary epithelial cells by immunohistochemistry, whereas 83% of the carcinoma cells showed positive staining for the LR (9). Importantly, overexpression of leptin was observed in 92% of breast tumors examined but in none of the normal breast epithelium (9). In recent years, many laboratories including our own have shown that leptin increases proliferation of different cancer cell types via multiple signaling pathways including Stat3/extracellular signal-regulated kinase/Akt signaling (10–13). The therapeutic potential of inhibition of leptin has been evaluated to some extent in diseases associated with metabolic syndrome, but the importance of inhibition of leptin signaling in carcinogenesis is still not known and is an active area of research.

Epidemiological results are compelling to show that dietary intake of cruciferous vegetables is protective against the risk of various types of malignancies including breast cancer (14–16). A case-control study involving >300 breast cancer patients and matched controls showed an inverse correlation between urinary levels of isothiocyanates (ITCs) as a biological measure of cruciferous vegetable intake and breast cancer risk (15). Similarly, another study showed that consumption of broccoli was inversely associated with premenopausal breast cancer risk (16). It is now widely accepted that anticarcinogenic effect of cruciferous vegetables is due to chemicals with an isothiocyanate functional group (N=C=S) that occur naturally as glucosinolates in a variety of edible cruciferous vegetables, including garden cress, broccoli, mustard, horseradish, watercress and cabbage (17). Benzyl isothiocyanate (BITC) is one of the best-studied members of the ITCs (18). Several studies have shown that broccoli is a rich source of BITC and isothiocyanates are much desired. We showed previously that benzyl isothiocyanate (BITC), a constituent of edible cruciferous vegetables (e.g. garden cress), confers significant protection against mammary carcinogenesis in a transgenic mouse model (23,24).

The present study was designed to explore if BITC functions as an antagonist of leptin. We specifically investigated the protective effect of BITC against the oncogenic actions of leptin. Intriguingly, we found that BITC inhibits the effect of leptin on malignant properties of breast cancer cells including migration and invasion and also inhibits important downstream molecules of leptin signaling.

Abbreviations: BITC, benzyl isothiocyanate; ChIP, Chromatin immunoprecipitation; ECIS, electric cell-substrate impedance sensing; ER, estrogen receptor; ITC, isothiocyanate; NF-kB, nuclear factor-kappaB; PCR, polymerase chain reaction; SFN, 1,3′-sulforaphane.

© The Author 2010. Published by Oxford University Press. All rights reserved. For Permissions, please email: journals.permissions@oup.com
Materials and methods

Cell culture and reagents

The human breast cancer cell lines, MCF-7 and MDA-MB-231, were obtained from the American Type Culture Collection and maintained in Dulbecco’s modified Eagle’s medium supplemented with 10% fetal bovine serum (Gemini Bioproducts, Woodland, CA) and 2 μM L-glutamine (Invitrogen, Carlsbad, CA). Cell line authentication was done by analysis of known genetic markers or response (e.g. expression of estrogen receptor and p53 and estrogen responsiveness). For treatment, cells were seeded at a density of 1 x 10^5/100 mm tissue culture dish. After 24 h of serum starvation, the culture media were changed to serum-free media containing treatments as indicated. Cultures were treated with human recombinant leptin (Sigma–Aldrich, St Louis, MO) at 100 ng/ml (12) and/or BITC (LKT Laboratories, St Paul, MN). Nα-sulfosulfonate (SUL) (purity 98.3%) was procured from LKT Laboratories. Antibodies for pS66 (Phospho-Stat3), Stat3, JAK2, pJAK2 (Phospho-JAK2), SRC1, p-Src (phospho-Src-Tyr416) and pSrc were purchased from Cell Signaling Technology (Danvers, MA). Antibodies against β-actin were purchased from Sigma–Aldrich.

Cell viability assay

Cells were seeded in 12-well plates at a density of 1 x 10^4 cells per well. After 12 h serum starvation, cells were pretreated with BITC (2.5 and 5 μM) or SUL (10 and 20 μM) for 2 h and then treated with leptin (100 ng/ml) for 22 h in the absence or presence of BITC. Effect of leptin and/or BITC treatments on cell viability was determined by trypan blue dye exclusion assay as described by us previously (25).

Clonogenicity assay

To perform colony formation assay (26), MCF-7 and MDA-MB-231 cells (single-cell suspension) were plated in 12-well plates at a density of 2500 cells per well overnight. The following day, cells were treated with leptin and/or BITC treatments as indicated and the medium was replaced with fresh medium containing treatment every 3 days. After a 10 day treatment period, the medium was removed and cells colonies were stained with crystal violet (0.1% in 20% methanol). Colony numbers were assessed visually and colonies containing >50 normal-appearing cells were counted. Pictures were taken using a digital camera. All experiments were performed at least three times in triplicate.

Anchorage-independent growth assay

Anchorage-independent growth of MCF-7 and MDA-MB-231 cells was determined by colony formation on soft agar (26). Briefly, equal volumes of agar (1.2%) and complete medium were mixed to make 0.6% agar growth medium solution in six-well tissue culture plates. Cells (2 x 10^3 cells per well) were suspended in media with or without leptin or BITC treatment as indicated followed by mixing with equal volume of agar (0.6%). Cell suspension agar mix (2 ml) was then added to each well. Plates were incubated at 37°C with 5% CO2 in humidified incubator for 3 weeks, and media with or without treatment were added every 3 days. Colonies were stained with 0.005% crystal violet in phosphate-buffered saline for 1 h at room temperature and observed using Olympus IX50 inverted microscope. Colonies were counted in five randomly selected fields at x10 magnification. Results are expressed as number of colonies counted. All experiments were performed three times in triplicate.

Migration assay

To perform migration assay (27), cells were plated into 24-well cell culture plates, pre-coated with human fibronectin (5 μg/cm2; Sigma–Aldrich). Cells were allowed to grow in 10% fetal bovine serum containing Dulbecco’s modified Eagle’s medium to confluence and then were washed with serum-free medium and serum starved for 16 h. A 1 mm wide scratch was made across the cell layer using a sterile pipette tip. After washing with serum-free medium twice, Dulbecco’s modified Eagle’s medium containing 10 μg/ml human fibronectin was added to replace matrix depleted with the cells. Plates were photographed immediately after scratching. Cells were treated with leptin and/or BITC as indicated. Plates were photographed after 24 and 48 h at the identical location of the initial image. All experiments were performed at least three times.

Electric cell-substrate impedance sensing wound-healing assay

Wound-healing assay was performed using the electric cell-substrate impedance sensing (ECIS) (Applied BioPhysics, Troy, NY) technology (27). For wound-healing assay, cells were grown to confluence on ECIS plates and serum starved for 16 h. The ECIS plates were submitted to an elevated voltage pulse of 40 kHz frequency, 3.5 V amplitude and 30 s duration, which led to the death and detachment of cells present on the small active electrode resulting in a wound normally healed by cells surrounding the small active electrode that have not been submitted to the elevated voltage pulse. Cells were immediately treated with leptin and/or BITC as indicated. Wound healing was then assessed by continuous resistance measurements for 24 h. All experiments were performed at least three times in triplicates.

Invasion assay

For an in vitro model system for metastasis, a Matrigel invasion assay (12) was performed using a Matrigel invasion chamber from BD Biocoat Cellware (San Jose, CA). Cells were seeded at a density of 1 x 10^5 cells per insert and cultured overnight. After 16 h of serum starvation, the culture media were changed to serum-free media containing treatments as indicated. Triplicate wells were used for each treatment. Cells were treated with leptin and/or BITC as indicated. After 24 h of incubation, cells remaining above the insert membrane were removed by gentle scraping with a sterile cotton swab. Cells that had invaded through the Matrigel to the bottom of the insert were fixed in methanol for 10 min. After washing with phosphate-buffered saline, the cells were stained with hematoxylin–eosin. The insert was subsequently washed with phosphate-buffered saline and briefly air-dried and mounted. The slides were coded to prevent counting bias, and the number of invaded cells on representative sections of each membrane were counted under light microscope. The number of invaded cells for each experimental sample represents the average of triplicate wells. All experiments were performed at least three times.

Western blotting

Whole cell lysate (12) was prepared by scraping MCF-7 and MDA-MB-231 cells in 250 μl of ice-cold modified RIPA buffer [50 mM Tris–HCl (pH 7.4), 150 mM NaCl, 1 mM ethylenediaminetetraacetic acid, 1% NP-40, 0.25% Na-deoxycholate, 1 mM phenylmethylsulfonyl fluoride, 10 μg/ml aprotinin, 10 μg/ml leupeptin, 1 mM Na3VO4, and 1 mM NaF]. The lysate was sonicated for 30 s, vortexed for 1 min, and incubated for 30 min on ice followed by centrifugation at 12 000 rpm at 4°C for 15 min to clear the cellular debris. Protein was quantified using the Bradford protein assay kit (Bio-Rad, Hercules, CA). Equal amount of lysate protein was resolved on sodium dodecyl sulfate–polyacrylamide gel, transferred to nitrocellulose membrane and western blot analysis was performed. Immunodetection was done by using enhanced chemiluminescence (ECL system, Amersham Pharmacia Biotech, Arlington Heights, IL) according to manufacturer’s instructions.

Immunohistochemical analysis for pSTAT3 in MDA-MB-231 xenografts

We have shown previously that BITC administration retards growth of MDA-MB-231 cells implanted in female athymic mice (23). We used tumor sections from the same study to determine the effect of BITC administration on expression of pSTAT3 by immunohistochemistry. Immunohistochemistry was performed essentially as described by us previously for other proteins (23,24). At least four non-overlapping representative images from each tumor section from five mice of each group were captured using ImagePro software for quantitation of pSTAT3 expression.

RNA interference and luciferase assay

Human cyclin D1 promoter reporter construct used for luciferase assay (~1745-CD1-Luc) was a kind gift from Dr Richard Pestell (Georgetown University, Washington, DC) (28). The plasmid expressing a constitutively active form of Stat3 (Stat3-FLAG) was a kind gift from Dr James E. Darnell (The Rockefeller University, New York, NY) (29). The nuclear factor-kappaB (NF-kB)-luciferase reporter plasmid was generously provided by Dr Aming Lin (University of Chicago, Chicago, IL). Cells were transfected with expression vectors and luciferase construct and treated with leptin and/or BITC as indicated using Lipofectamine 2000 (Invitrogen) according to the manufacturer’s instructions. Forty-eight hours post-transfection, the cells were harvested and luciferase and Renilla activities were measured using a dual luciferase kit (Promega, Madison, WI). The relative firefly luciferase activities were calculated by normalizing transfection efficiency according to Renilla luciferase activities. For RNA interference, cells were seeded in six-well plates and transfected at 50% confluency with 100 nM of control siRNA or Stat3-targeted siRNA (Invitrogen) using Oligofectamine according to the manufacturer’s instructions. Twenty-four hours post-transfection, the cells were harvested and luciferase and Renilla activities were measured using a dual luciferase kit (Promega). The relative firefly luciferase activities were calculated by normalizing transfection efficiency according to Renilla luciferase activities. The experiments were performed three times in triplicate.

Chromatin immunoprecipitation

Chromatin immunoprecipitation (ChIP) analyses were performed using our published procedure (12) with following modifications. Chromatin samples were prepared using the protocol described in the chromatin immunoprecipitation (ChIP) kit (Upstate) according to manufacturer’s instructions. Chromatin-DNA complexes were prepared using 5 μg of chromatin per sample. The samples were incubated with the appropriate antibodies and then immunoprecipitated with protein A-Sepharose beads. The supernatants were analyzed by quantitative real-time PCR using the specific primers for the human cyclin D1 promoter region.
were sonicated on ice three times for 10 s each (i.e. until the average length of sheared genomic DNA was 1–1.5 kb) followed by centrifugation for 10 min. The immunoprecipitated DNA was ethanol precipitated and resuspended in 25 μl of water. Total input samples were resuspended in 100 μl of water and diluted 1:100 before polymerase chain reaction (PCR) analysis. Initially, PCR was performed with different numbers of cycles and/or dilutions of input DNA to determine the linear range of amplification; all results shown fall within this range. Following 28–30 cycles of amplification, PCR products were run on 1% agarose gel and analyzed by ethidium bromide staining. All ChIP assays were performed at least thrice.

Results

**BITC-treatment inhibited oncogenic actions of leptin in human breast cancer cells**

We showed previously that leptin increases proliferation and growth of breast cancer cells via activation of multiple downstream signaling pathways including Stat3 (10–12,27). Because BITC has been shown to inhibit Stat3, we postulated that the oncogenic effects of leptin may be reversed in the presence of BITC. Here, we experimentally tested this hypothesis using well-characterized human breast cancer cell lines (MCF-7 and MDA-MB-231) as a model. The effect of BITC on cell viability of breast cancer cells was determined by trypan blue exclusion assay and the results are summarized in Figure 1A. Leptin treatment significantly stimulated survival of MCF-7 and MDA-MB-231 cells, whereas the survival of both cell lines was decreased significantly after exposure to BITC in a concentration-dependent manner. Cell viability was reduced to ~50–60% with 2.5 μM BITC treatment, whereas higher concentration (5 μM) was more inhibitory. The MCF7 cells were relatively more sensitive to the leptin-induced cell proliferation as compared with MDA-MB-231 cells. In addition, BITC conferred significant protection against leptin-stimulated cell viability of MDA-MB-231 and MCF-7 cells (Figure 1A). Breast cancer cells were treated with various concentrations of BITC and subjected to clonogenicity and anchorage-independent growth assay. Dose dependent and statistically significant inhibition of clonogenicity and soft-agar colony formation was observed in the presence of BITC (supplementary Figure S1 is available at Carcinogenesis Online). Consistent with these observations, BITC (2.5 μM) treatment not only reduced leptin-induced clonogenicity (Figure 1B) but also anchorage-independent growth in both cell lines (Figure 1C).

Cancer progression is a multistep process that involves invasion of basement membrane by tumor cells and migration to points far from a given primary tumor mass leading to metastasis (30). We examined the effects of leptin and/or BITC treatments on invasion and migration properties of breast carcinoma cells using Matrigel invasion and scratch migration assay. BITC treatment inhibited migration and invasion potential of breast cancer cells in a concentration-dependent manner (supplementary Figure S2 is available at Carcinogenesis Online). As expected, we found that leptin increased migration of breast carcinoma cells, whereas BITC inhibited migration in a conventional scratch migration assay. Importantly, BITC treatment also inhibited migration of MCF-7 and MDA-MB-231 breast cancer cells in the presence of leptin overcoming its strong pro-migratory potential (Figure 2A). For quantitative determination of alteration in migration potential of breast cancer cells upon treatment with leptin and/or BITC, we performed a quantitative real-time impedance assay using an ECIS-based technique (27). As expected, confluent cells showed high resistance values. Confluent cells were subjected to high voltage pulse that resulted in drop in resistance indicating death and detachment of cells present on the small active electrode followed by leptin and/or BITC treatments. Leptin treatment resulted in higher resistance showing increased migration of cells surrounding the small active electrode that were not submitted to the elevated voltage pulse. BITC treatment inhibited migration resulting in lower resistance values. BITC treatment also inhibited leptin-induced migration in combination treatment (Figure 2B). Next, we performed Matrigel invasion assay to examine the effect of BITC on leptin-induced invasion potential of breast carcinoma cells. As evident from Figure 2C, treatment increased invasion of cancer cells through Matrigel in comparison with untreated cells, whereas BITC treatment inhibited invasion of breast cancer (MCF-7 and MDA-MB-231) cells. Interestingly, we found that BITC treatment efficiently inhibited leptin-induced increased invasion of cancer cells (Figure 2C). Collectively, these results show that BITC treatment can effectively inhibit leptin-induced clonogenicity, anchorage-independent colony formation, migration and invasion of breast carcinoma cells.

**BITC treatment suppressed leptin-induced Stat3 phosphorylation in breast cancer cells**

Binding of leptin to LR (Ob-Rb) phosphorylates conserved tyrosine residues and these phosphorylation events are important for subsequent signaling events including JAK and Stat3 activation (12). Our previous studies have shown direct involvement of JAK/Stat3,
phosphatidylinositol 3-kinase/Akt and extracellular signal-regulated kinase signaling in pro-cancerous actions of leptin (12). To investigate hierarchy of these events, we previously found that leptin signaling is transmitted mainly by the JAK/Stat3 pathway as inhibition of Stat3 activation further inhibits leptin-induced phosphatidylinositol 3-kinase/Akt and extracellular signal-regulated kinase signaling as well as leptin-induced proliferation (10,11). These studies put forth Stat3 activation as an integral event in leptin signaling. We sought to determine the underlying molecular mechanism by which BITC treatment inhibited oncogenic actions of leptin. MCF-7 and MDA-MB-231 cells were treated with leptin for 30 min or 1 h. Consistent with previous observations (12), Tyr705 phosphorylation of Stat3 was increased markedly in leptin-treated MCF-7 and MDA-MB-231 cells, which was not due to an increase in protein level of total Stat3 (Figure 3A).

Fig. 2. BITC inhibits leptin-induced migration and invasion of breast carcinoma cells. (A) Breast cancer cells (MCF-7 and MDA-MB-231) were subjected to scratch migration assay. Culture media were replaced with media containing 100 ng/ml leptin, 2.5 μM BITC alone and in combination or untreated media (U). The plates were photographed at the identical location of the initial image (0 h) at 24 h. The results shown are representative of three independent experiments performed in triplicates. The histogram shows the fold change in migration. *P<0.01 compared with untreated controls; **P<0.001 compared with untreated cells; ***P<0.005 compared with leptin treatment. All the experiments were performed thrice in triplicates. (B) MCF-7 and MDA-MB-231 cells were grown to confluence in ECIS plates, serum starved for 16 h and subjected to an elevated voltage pulse of 40 kHz frequency at 3.5 V amplitude for 30 s to incite a wound. The cells were immediately treated with leptin and/or BITC as indicated. The wound was then allowed to heal from cells surrounding the small active electrode that did not undergo the elevated voltage pulse. Resistance was measured before and after the elevated voltage pulse application as described in the ‘Materials and Methods’. The measurements were stopped 24 h after the creation of wound. All the experiments were performed thrice in triplicates. (C) MCF-7 and MDA-MB-231 cells were cultured in Matrigel invasion chambers followed by treatment with leptin, BITC alone and in combination for 24 h. The number of cells that invaded through the Matrigel was counted in five different regions. The slides were blinded to remove counting bias. The results show mean of three independent experiments performed in triplicates. *P<0.005 compared with untreated controls; **P<0.001 compared with untreated controls; ***P<0.001 compared with leptin-treated cells.
time-dependent manner. For example, the level of pStat3Tyr705 was reduced by 70% by a 24 h exposure to 2.5 and 5 μM BITC (Figure 3B). Phosphorylation of Stat3 is mediated by various kinases, including Src and JAK2 (33,34). We examined the level of pJAK2 to determine if BITC-mediated inhibition of Stat3 activation was due to suppression of JAK2 phosphorylation. The level of pJAK2 was also reduced in a concentration- and time-dependent manner upon BITC treatment (Figure 3B). BITC-mediated suppression of JAK2 phosphorylation correlated with a marked decrease in its protein level (Figure 3B). We also examined the level of pSrc to determine its involvement in BITC-mediated inhibition of Stat3 activation. BITC treatment reduced the level of pSrc in a time-dependent manner (Figure 3B). In a recent study, we showed that BITC administration retarded growth of MDA-MB-231 cells implanted in female athymic mice (23). We used tumor sections from the same study to determine effect of BITC administration on expression of pSTAT3 protein by immunohistochemistry. We found decreased phosphorylation of Stat3 in tumors from BITC-treated mice in comparison with control tumors (Figure 3C).

Stat3 was a critical target of BITC-mediated inhibition of leptin signaling

We have shown previously that one of the targets of leptin action is cyclin D1, and leptin treatment induces upregulation and transactivation of cyclin D1 (12). Next, we investigated if BITC treatment inhibits leptin-induced cyclin D1 transactivation using a cyclin D1 promoter luciferase construct, -1745-CD1-Luc (Figure 4A). The MCF-7 and MDA-MB-231 cells transfected with -1745-CD1-Luc demonstrated a 4- to 6-fold increase in luciferase activity in response to leptin exposure. BITC treatment alone significantly inhibited cyclin D1 transactivation indicating that BITC negatively regulates the activity of the cyclin D1 promoter. Interestingly, we also found that the leptin-stimulated cyclin D1 transactivation was partially but
cytokine action mediated by Stats in other systems. We further sought
to determine if BITC treatment can inhibit direct participation of Stat3 in
leptin-mediated cyclin D1 gene regulation using ChIP assay. Using
specific antibodies against Stat3, formaldehyde cross-linked protein–
chromatin complexes were immunoprecipitated from MCF-7 and
MDA-MB-231 cells cultured with or without leptin and/or BITC.
Resulting precipitated genomic DNA was then analyzed by PCR using
primers spanning the Stat3-binding elements in the promoter region of
the cyclin D1 gene. As shown in Figure 4B, ChIP analysis with anti-
Stat3 antibody revealed that Stat3 was associated with the cyclin
D1 promoter in the presence of leptin, whereas BITC treatment ab-
rogated leptin-induced recruitment of Stat3 on cyclin D1 promoter
(Figure 4B). Previously, we have shown that leptin-induced Stat3
recruits histone acetyltransferase activity containing coactivator such
as SRC-1 to cyclin D1 promoter. We analyzed the recruitment of
SRC1 to cyclin D1 promoter in the presence of BITC. We found that
BITC inhibits recruitment of SRC1 to cyclin D1 promoter
(Figure 4B).

BITC treatment inhibited leptin-induced recruitment of Stat3 to
cyclin D1 promoter indicating that Stat3 may be a pivotal point in
BITC-leptin cross talk. Therefore, the effect of overexpression of
constitutively active Stat3 on BITC-mediated inhibition of leptin sig-
ning was examined. Leptin-induced luciferase activity driven by the
cyclin D1 promoter was augmented by co-transfection of constituti-
tively active Stat3 (Stat3C). BITC treatment inhibited cyclin D1 trans-
activation with or without leptin treatment. Importantly, Stat3C
overexpression interfered with BITC-mediated inhibition of cyclin
D1 transactivation in the presence of leptin (Figure 5A). For further
confirmation, we proceeded to determine the effect of Stat3 protein
knockdown on BITC-mediated cyclin D1 transactivation in MCF-7
and MDA-MB-231 cells. The level of Stat3 and p-Stat3 protein was
reduced by 80% on transient transfection of breast cancer cells with
Stat3-targeted siRNA in comparison with cells transfected with a con-
trol non-specific siRNA (Figure 5B). Knockdown of Stat3 resulted in
a decrease in leptin-induced cyclin D1 transactivation. On the other
hand, Stat3 protein inhibition significantly potentiated inhibitory ef-
fects of BITC on cyclin D1 transactivation. We also observed a statis-
tically significant increase in BITC-mediated inhibition of cyclin D1
transactivation in the presence of leptin (Figure 5C). Collectively,
the results showed the involvement of Stat3 in BITC-leptin cross talk.
BITC treatment inhibited leptin-induced stat3 activation and recruit-
ment to cyclin D1 promoter resulting in inhibition of cyclin D1 trans-
activation in breast cancer cells.

Next, we questioned if inhibition of oncogenic effects of leptin
was unique to BITC. We addressed this question using SFN and MDA-
MB-231 cells. SFN, a synthetic racemic analogue of broccoli-derived
t-isomer, inhibits proliferation and induces apoptosis in many cancer
cell types (39). Similar to BITC, SFN treatment reduced cell viability
in the absence or presence of leptin stimulation (Figure 6A). In addi-
tion, SFN treatment inhibited invasion potential of breast cancer cells
(Figure 6B). SFN was also effective in inhibiting leptin-induced cell
invasion (Figure 6C).

Discussion
Adipose tissue is an active endocrine organ that secretes various
biologically active adipocytokines and provides a potential molecular
mechanism to link obesity with carcinogenesis (40). One of the im-
portant adipocytokine is leptin. Genetically, obese leptin-deficient
MMTV-TGF-α/Lepob and leptin-receptor-deficient MMTV-
TGF-α/Leprb female mice do not develop mammary tumors
providing additional evidence for the integral role of leptin and its
cognate receptor in mammary tumorigenesis (41,42). Analysis of
clinical samples showed that LRs were not detectable in normal mam-
mary epithelial cells by immunohistochemistry, whereas carcinoma
cells showed positive staining for OB-R (leptin receptor) in 83% of
the cases (9). Leptin has been characterized as a potent growth-stim-
ulating factor for breast cancer that also induces breast cancer cell
migration and invasion (27). Previous studies have shown that leptin

statistically significantly suppressed in the presence of BITC (Figure 4A).
Thus, BITC efficiently overcame the stimulatory effects of leptin
on cyclin D1 transactivation (Figure 4A). Cyclin D1 is also regulated
by NF-κB (35). We investigated the effects of BITC on NF-κB in
a luciferase assay. We found that BITC does not affect NF-κB activity
(supplementary Figure S3 is available at Carcinogenesis Online).

Gene expression analyses in response to cytokine signaling have
shown that Stat proteins are signaling molecules with dual functions
(36). Stat proteins not only transmit a signal from the cell surface to
the nucleus but also directly participate in gene regulation (36).
Cyclin D1 promoter contains binding sites for a number of transcription
factors (37,38). The consensus Stat-binding motifs (TTCNNNGAA)
at -247 and -481 were of particular interest for these studies because
these consensus sites have previously been shown to be sites for

Fig. 4. BITC inhibits leptin-induced recruitment of Stat3 to cyclin D1 promoter. (A) Schematic representation of cyclin D1 promoter showing two GAS consensus sites located at -481 and -247. MCF-7 and MDA-MB-231 cells were transiently transfected with the human cyclin D1 construct (-1745-
CD1-Luc) together with the internal control Renilla luciferase reporter and
and treated with leptin and/or BITC. Untreated cells are denoted with the letter ‘U’. Cell lysates were assayed for luciferase activity and activities were
normalized against the internal control. Data represents the mean of three
experiments. *P < 0.005 compared with untreated controls; **P < 0.01
compared with leptin-treated cells; #P < 0.001 compared with leptin treated cells. (B) Soluble chromatin was prepared from MCF-7 and MDA-MB-231
 cells, treated with leptin and/or BITC and immunoprecipitated with 5 μg of
specific antibody against pStat3 and SRC1 overnight at 4°C. The immune
complexes were pulled-down with protein A agarose/salmon sperm DNA
beads and washed extensively and cross-linking was reversed. The purified
dNA was analyzed by PCR using primers spanning the GAS sites (-481
and -247) at the cyclin D1 promoter. ChIP assay shows that BITC inhibits
leptin-induced Stat3 and SRC1 recruitment to cyclin D1 promoter.
induces proliferative responses both in ER-negative and ER-positive breast cancer cells (43,44). However, a recent study reported differential sensitivity to leptin treatment in breast cancer cells based on ER status (45). Discrepancy might be due to clonal variations of the cell lines used. We consistently found leptin-induced proliferation in both MCF-7 and MDA-MB-231 cells.

Reagents blocking leptin activity might prove useful for breast cancer patients with elevated leptin levels. Such neutralization can be achieved with soluble LRs that bind free leptin in the circulation, leptin antagonists that bind to, but do not activate LRs, and either specific anti-LR monoclonal antibodies (anti-LR mAbs) that bind to the receptor preventing leptin signaling or anti-leptin antibodies. Anti-LR mAbs have high molecular mass that ensures a long half-life in the circulation and good affinity for the receptor. But these mouse-generated mAbs need to be humanized to eliminate their potential immunogenicity. Nanobodies represent another approach to target LR and block the ligand-induced conformational change without interfering with the leptin-LR interaction. These recombinant,
monomeric small nanobodies are stable and easy to manipulate (46). They can selectively inhibit peripheral activity of leptin as they do not cross blood–brain barrier. Importantly, recent development of leptin muteins with antagonistic properties and other proteins blocking leptin activity present new possibilities for research. To date, all these agents to counteract leptin signaling are in initial stages of development. Therefore, there is a clear and compelling need to identify novel agents that are relatively safe but could suppress leptin signaling and its biological effects.

BITC, a promising cancer chemopreventive agent, has been shown previously to noticeably suppress the incidence and/or burden of mammary hyperplasia and carcinoma in female MMTV-neu mice (24). BITC-mediated prevention of mammary carcinogenesis in MMTV-neu correlates not only with reduction of cell proliferation and increased apoptosis but also infiltration of T cells (24). BITC has also been shown to impede MDA-MB-231 breast cancer xenograft growth in association with decreased tumor cell proliferation (23). Moreover, BITC treatment reduced migratory potential of MDA-MB-231 cells (23). On the other hand, leptin has been shown to increase tumor growth in vivo and augment proliferation, invasion and migration of breast cancer cells (27,47,48). The present study shows that BITC inhibits leptin-induced breast cancer growth and metastatic properties. Specifically, BITC treatment inhibits leptin-induced cloningogenicity, anchor-independent three dimensional colony formation, invasion and migration of breast carcinoma cells. Higher levels of leptin in obese breast cancer patients have been associated with higher risk for lymph node metastasis, larger tumor burden when compared with non-obese breast cancer patients. Our studies indicate that BITC treatment may be able to inhibit tumor metastasis owing to high leptin levels.

Examination of molecular pathogenesis of cancer has shown that disruption of the cell cycle in human tumors is a critical interface between hormonal signaling and tumorigenesis. The dysregulation of cell cycle control in cancer is compressed into a model where disruption of two parallel pathways is required for the occurrence of cancer. These two parallel pathways include either the p16 (INK4a) pathway or the p19 (ARF) pathway. Our studies show that leptin led to overexpression of cyclin D1 thus inactivating one arm of the signaling pathway leading to unchecked cell cycle progression and possibly tumorigenic growth. Cyclin D1 transgenic animals show perturbed mammary gland development with increased proliferation and previous lobular development (49). Tumors appear in 75% of mice but after a long latency compared with oncogenes c-myc, Ha-ras and c-neu, which induce tumors relatively early (49). The ability of cyclin D1 overexpression to induce mammary carcinoma and the necessity for cyclin D1 function for cell cycle progression raises the question of whether cyclin D1 is necessary for tumor development. So far, the relationship between overexpression of cyclin D1 and breast cancer outcome has been controversial with studies reporting both positive and negative findings (50,51).

We found that BITC treatment inhibited leptin-induced cyclin D1 transactivation.

Investigating the signaling pathways involved in leptin-induced cyclin D1 transactivation, we found that activation of JAK/Stat pathway regulates the transactivation of cyclin D1 gene (12). In a parallel study, Leslie et al. (52) reported that cyclin D1 is transcriptionally regulated by Stat3 and is required for transformation. We previously showed that leptin-activated-Stat3 binds to its cognate binding sites in cyclin D1 promoter, acting as a transcription factor, leading to hyperacetylation and overexpression of cyclin D1 gene. Inhibition of JAK/Stat pathway significantly reduced leptin-induced growth of breast carcinoma cells (12). Here, we show that BITC inhibits leptin-induced Stat3 activation by phosphorylation at Tyr705. Our study shows that BITC-mediated inhibition of Stat3 activation in breast cancer cells is accompanied by suppression of JAK2 activation. The kinetics of BITC-mediated inhibition of JAK2 is similar to that of Stat3 inhibition. BITC treatment also reduces protein levels of JAK2. Stat3 activation leads to dimerization, nuclear translocation and recognition of Stat3-specific DNA-binding elements and activation of target gene transcription (33,34). We found that BITC treatment inhibits leptin-induced Stat3 recruitment to cyclin D1 promoter. Furthermore, Stat3 activation represents a pivotal point in leptin-BITC cross talk as overexpression of constitutively active Stat3 interferes with BITC-mediated inhibition of leptin action, whereas Stat3 knockdown potentiates BITC-mediated inhibition of leptin action.

In summary, the present study provides concrete experimental evidence in support of a novel function of BITC as a potential antagonistic agent against oncogenic actions of leptin. Considering the high prevalence of obesity in the developed world, our study has the potential to significantly impact the vast majority of obese breast carcinoma patients by providing new possibilities for research and ultimately therapy using BITC for inhibiting invasion and migration properties of tumor cells and improving overall prognosis.

**Supplementary material**

Supplementary Figures S1–S3 can be found at http://carcin.oxfordjournals.org/

**Funding**

United States Public Health Service (NCI-R01 CA129347-04, NCI-R01 CA142604-01 to S.V.S.; NIDDK-K01 DK077137, NIDDK-R03 DK089130 to N.K.S.; NCI-R01 CA131294 to D.S.).

**Conflict of Interest Statement:** None declared.

**References**


Received July 21, 2010; revised December 3, 2010; accepted December 8, 2010