Effect of curcumin on irradiated and estrogentransformed human breast cell lines

GLORIA M. CALAF^{1,2}, CARLOS ECHIBURÚ-CHAU¹, GENGYUN WEN², ADAYABALAM S. BALAJEE² and DEBASISH ROY³

¹Instituto de Alta Investigación, Universidad de Tarapacá, Arica, Chile; ²Center for Radiological Research, Columbia University Medical Center, New York, NY, USA; ³Department of Natural Science, Hostos College of the City University of New York, New York, NY, USA

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Abstract. Curcumin (diferuloyl methane) is a well known antioxidant that exerts antiproliferative and apoptotic effects. Curcumin effect was evaluated in a breast cancer model that was developed using the immortalized breast epithelial cell line MCF-10F after exposure to low doses of high LET (linear energy transfer) α particles (150 keV/ μ m) of radiation, and subsequently cultured in the presence of 17\beta-estradiol (estrogen). This model consisted of human breast epithelial cells in different stages of transformation: i) MCF-10F; ii) Estrogen cell line; iii) a malignant Alpha3 cell line; iv) a malignant and tumorigenic, Alpha5 cell line; and v) a cell line derived from Alpha5 injected into the nude mice that gave rise to Tumor2 cell line. Curcumin decreased anchorage-independent growh in transformed breast cancer cell lines in comparison to their counterparts and increased the percentage of cells from G₀/G₁ with a concomitant increase in G₂/M phases, as well as a decrease in PCNA and Rho-A protein expression. Among the oncogenes, c-Ha-Ras and Ras homologous A (Rho-A) are important cell signaling factors for malignant transformation and to reach their active GTP bound state, Ras proteins must first release bound GDP mediated by a guanine nucleotide releasing factor (GRF). Then curcumin decrease RasGRF1 protein expression in malignant cell lines. Further, differential expression levels of cleaved (ADP) ribose polymerase 1 (PARP-1) and phosphorylated histone H2AX (y-H2AX) were observed after curcumin treatment. It seems that PARP-1 similar to H2AX, confers cellular protection against radiation and estrogen-induced DNA damage mediated by curcumin. Therefore, targeting either PARP-1 or H2AX may provide an effective way of maximizing the therapeutic value of antioxidants for cancer prevention.

E-mail: gmc24@columbia.edu

Key words: curcumin, c-Ha-Ras, breast cells, radiation, estrogen

Introduction

Among the antioxidants, curcumin (diferuloyl methane), is a well-known major dietary natural yellow pigment derived from the rhizomes of turmeric the herb knowns as *Curcuma longa* originary from India and South Asia. It possesses antiproliferative, anticarcinogenic and chemopreventive potential (1,2). The antitumor efficacy of curcumin has been documented in animal models as well as in phase I clinical trials with multiple cancer types (3-7). The molecular targets and therapeutics of curcumin in health and disease have been studied before (5-7). These studies have reported that curcumin had biological and medicinal properties as well as anticancer potential in preclinical and clinical studies. Numerous studies in animals have demonstrated that curcumin has potent chemopreventive activity against a wide variety of tumors. Curcumin has great potential in the prevention and treatment of cancer.

Curcumin has been shown to be chemopreventive and chemotherapeutic by blocking tumor initiation (8-13) induced by benzo[a]pyrene and 7,12 dimethylbenz[a]anthracene; it suppressed phorphol ester-induced tumor promotion; suppressed carcinogenesis of the skin, forestomach; and colon in mice. It has been shown to prevent cancer in colon, skin stomach, duodenum following oral administration (7-9).

Curcumin inhibited cell proliferation, invasion, angiogenesis and metastasis of different cancers through interaction with multiple cell signaling proteins (1). Various molecular targets modulated by this agent include transcription factors, growth factors and their receptors, cytokines, enzymes, and genes regulating cell proliferation and apoptosis (14-17). Curcumin inhibits activation of transcriptional factors such as (AP)-1, nuclear factor-kappa B (NF- κ B) and many others that are up-regulated in most cancers. In normal primary human cells, curcumin either stimulates or inhibits proliferation, but does not appear to induce apoptosis. Curcumin reversibly arrests normal mammary epitelial cells at G₀/G₁ phases. However, apoptosis by curcumin was reported in mammary epithelial carcinoma cells at the G₂ phase of cell cycle (14-19).

Among the oncogenes, *c-Ha-Ras* and *Ras homologous A* (*Rho-A*) from the Ras super family have been shown to promote both cell proliferation and invasion indicating their importance in malignant transformation (20-31). A critical step in the

Correspondence to: Dr Gloria M. Calaf, Instituto de Alta Investigación, Universidad de Tarapacá, Calle Antofagasta no. 1520, Arica, Chile

stimulation of cell surface receptors by their ligand involves the accumulation of Ras proteins in their active GTP-bound state. To reach their active GTP bound state, Ras proteins must first release bound GDP mediated by a guanine nucleotide releasing factor (GRF). Members of Rho family proteins sometimes act downstream of Ras. The Rho GTPases have been identified as regulators of cytoskeletal reorganization in addition to their effects on cell growth and it has been shown that they are over-expressed in human tumors (20-22).

The DNA damage response is a complex process involving multiple DNA repair, cell survival, and cell death pathways with damage specificity to different types of DNA damage (34-37). The DNA damage includes double-strand breaks (DSB), single strand breaks, base damage, bulky adducts, intra/interstrand cross links, and breakdown of replication fork lesions where one of the key proteins in the base excision repair pathway is the Poly adenosine diphosphate ribose polymerase-1 (PARP-1) (6,19,32-34). PARP-1 is a 116-kDa nuclear protein that appears to be involved in DNA repair predominantly in response to environmental stress (32,33). It is important for cells to maintain their viability and cleavage of PARP facilitates cellular disassembly and serves as a marker of cells undergoing apoptosis.

Accumulation of phosphorylated histone H2AX, also called γ -H2AX, is a marker for DSB (35-37). The H2AX, member of the family H2A, is a subunit of the nucleosome. The phosphorylation of this protein at the serine 139 in the C-tail serine-glutamine-glutamate motif is one of the earliest responses of mammalian cells to ionizing radiation-induced DNA double-stand breaks (DSB) (35-37). H2AX forms discrete foci at the sites of DSBs, facilitates the remodeling complexes to the sites of DNA damage, and influences both the efficiency and fidelity of DSB repair.

To gain insights into the effects of curcumin on breast carcinogenesis an established *in vitro* experimental breast cancer model (Alpha model) (38-40) induced by radiation and estrogen was used. The aim of this work was to evaluate the effect of curcumin in human breast epithelial cells transformed by the effect of radiation in the presence of estrogen and to identify the biological processes that are altered during the malignant transformation of normal breast epithelial cells.

Materials and methods

Experimental breast cancer model. The spontaneously immortalized breast epithelial cell line, MCF-10F (ATCC, Manassas, VA) retains all the characteristics of normal epithelium in vitro, including anchorage-dependence, non-invasiveness and nontumorigenicity in nude mice. This cell line was exposed to low doses of high LET (linear energy transfer) alpha particles (150 keV/ μ m) in the presence of estrogen (17 β -estradiol) (38). This model consisted of human breast epithelial cells in different stages of transformation: i) normal cells, MCF-10F; ii) MCF-10F cell line treated with estrogen (10⁻⁸ M), called Estrogen cell line; iii) a malignant non-tumorigenic, Alpha3 cell line; iv) a malignant and tumorigenic, Alpha5 cell line; and v) a malignant and tumorigenic cell line derived from Alpha5 injected into the nude mice giving rise to a tumor from which a cell line named Tumor2 was originated. The cell lines were cultured with Dulbecco's modified Eagle's media (DMEM)/F-12 (1:1) supplemented with 100 U/ml penicillin, 100 mg/ml streptomycin, 2.5 mg/ml amphotericin B (all from Life Technologies, Grand Island, NY), 10 mg/ml insulin (Sigma-Aldrich, St. Louis, MO), 5% equine serum (Biofluids, Rockville, MD), 0.5 mg/ml hydrocortisone (Sigma) and 0.02 mg/ml epidermal growth factor (Collaborative Research, Bedford, MA). The cells were incubated at 37°C with 5% CO₂ up to 70% confluence.

Growth rate of cells. The growth rate was determined by the doubling time of the cells seeded onto a flask T25 ($1x10^5$ cells per flask at initial time). To determine the population doubling time cells were counted 24, 48, and 72 h after plating that was calculated from the exponential portion of the growth curve. Dose-response curve of curcumin was analyzed at doses from 10 to 60 μ M after 48-h treatments.

Anchorage-independent assay. To test for cell growth in semisolid medium, curcumin-treated cells were trypsinized and plated at a density of 1×10^5 cells in 10 ml of 0.35% agarose over a 0.7% agar base in 100-mm culture dishes as described (23,24). Media were replenished every day and colonies were scored after 21 days in culture.

Cell cycle analysis. The cells were trypsinized then fixed in 70% ethanol and washed with cold PBS. To assess the DNA content the cells were incubated for 30 min with propidium iodide staining that was used for measuring by fluorescence-activated cell sorting (FACS propidium iodide/RNase staining buffer) (BD Pharmingen[™]), covered with aluminium foil and measured using flow cytometry. Percentage of cells in each phase of the cell cycle was analyzed using BD CellQuest Pro software (Becton-Dickinson, Co.) with an ultraviolet excitation laser beam.

Western blot analysis. Cells were lysated with 1 ml lysis buffer (pH 7.2) [Tris-Base (50 mM), EDTA (1 mM), NaCl (100 mM), PMSF (1 mM), Orthovanadate (1 mM), Triton X-100 (0.1%)] and centrifuged (10000 rpm x 10 min). Cellular proteins from the supernatant were dissolved in SDS-PAGE sample solution [60 mM Tris, pH 6.5, 10% (w/v) glycerol, 5% (w/v) β-mercaptoethanol, 20% (w/v) SDS, and 0.025% (w/v) bromophenol blue] and denatured by boiling (100°C x 5 min). The total amount of protein used was 20 μ g per lane with standard protein markers (Bio-Rad Laboratories, Hercules, CA). After fractionation by SDS-PAGE on gels (7x14 cm), proteins were electroblotted onto nitrocellulose membrane (Amersham, Biosciences, UK) using a blotting apparatus (Bio-Rad Laboratories, Hercules, CA). Prestained SDS-PAGE (Standards) blots were blocked for 2 h in 10% defatted dry milk-TBS-0.1% Tween and then incubated for 2 h at room temperature with corresponding primary antibodies. The following antibodies were used: PCNA (PC11, sc53407); Ras-GRF1(C18, sc863); RhoA (26C4, sc418) (all from Santa Cruz Biotechnology, CA), cleaved PARP (46D11, rabbit mAb) and phospho-H2AX (Ser 139, 20E3) (both from Cell Signaling, CA) (1: 200) followed by incubation with secondary peroxidaseconjugated monkey, mouse or anti-rabbit IgG (1:5000) (Cell Signaling, CA) in 5% defatted dry milk-TBS-0.1% Tween. All steps were performed at room temperature, and blots were rinsed between incubation steps with TBS-0.1% Tween. Cell blots were probed with mouse anti-actin antibody as control. Immunoreactive bands were visualized by using the ECLTM detection reagent (Amersham, Dübendorf, Switzerland) and



Figure 1. Effect of curcumin (dose-response curve of 0-60 μ M) on the percentage of MCF-10F, Estrogen, Alpha3, Alpha5 and Tumor2 cell lines after 2-day treatment. It was determined by using the Coulter Counter apparatus.

exposure of the membrane to X-ray film. Protein levels were determined with the Bicinchoninic acid method (Bio-Rad) and bovine serum albumin to perform the standard curve.

Results

Dose response curve of curcumin. To determine the dose to be used in all the experiments 0-60 μ M of curcumin was used for 48 h of treatment with the five cell lines. Results in Fig. 1 show that the mean LD₅₀ was at 25 μ M with the exception of Alpha3 cell line which showed an LD₅₀ value of 15 μ M. Then, all the experiments were carried out with either 15 or 30 μ M curcumin.

Anchorage-independent growth. Previous results with these five cell lines indicated that Alpha3, Alpha5 and Tumor2 were able to form colonies in soft agar. Curcumin at 30 μ M concentration decreased the colony formation under similar conditions in Alpha3, Alpha5 and Tumor2 cell lines (Fig. 2A). In contrast, there was no colony formation in the control and Estrogen cell lines with or without curcumin treatment.

Effect of curcumin on the cell cycle. Effect of curcumin on cell cycle profiles of the five cell lines with (30 μ M) and without curcumin treatment was studied (Fig. 2B and Table I) Results indicated that G₁, S and G₂/M phases were altered by curcumin by reducing the percentage of cells in G₀/G₁ phase and by increasing the proportion of G₂/M phase cells. The effect was more pronounced in Alpha3 and Alpha5 than in Tumor2 cell line.

Effect of curcumin on cell proliferation. We next investigated the level of the major DNA replicating factor, proliferating cell nuclear antigen (PCNA). Immunoperoxidase staining revealed that this cell cycle modulator was reduced with 15 μ M curcumin in the malignant cell lines Alpha3, Alpha5 and Tumor2. In contrast, MCF-10F and Estrogen cell lines did not show any change (Fig. 3A).

Table I. Effect of curcumin on cell cycle of breast cancer epithelial cell lines.

Cell lines	$G_0/G_1 (\%)^a$	S (%) ^a	$G_2/M~(\%)^a$
MCF-10F	31.85	11.48	22.88
MCF-10F + Cur	33.39	16.63	19.76
Estrogen	15.66	6.42	21.88
Estrogen + Cur	17.92	10.14	18.47
Alpha3	42.44	8.04	19.15
Alpha3 + Cur	33.90	12.81	29.12
Alpha5	52.96	7.22	22.02
Alpha5 + Cur	30.5	14.57	30.77
Tumor2	14.03	8.41	19.11
Tumor2 + Cur	20.42	7.88	26.51

+ Cur, dose of 30 µM curcumin. ^aPercentage of cells.

Effect of curcumin on Ras activated state. To analyze the Ras active GTP bound state, the RasGRF1 protein expression was assessed. Expression and quantification of effect of 15 μ M curcumin in all the cell lines are shown in Figs. 3B and 4. Results indicated that curcumin decreased RasGRF1 protein expression in the control MCF-10F, Apha5 and Tumor2 cell lines by fluorescence staining intensities (Fig. 3B). RasGRF1 protein expression increased in control and Estrogen cell line by the effect of curcumin but decreased in Alpha3, Apha5 and Tumor2 cell lines in the presence of curcumin (Fig. 4).

Effect of curcumin on regulation of actin cytoskeleton. RhoA is a small GTPase protein known to regulate the actin cytoskeleton in the formation of stress fibers and it is generally distributed in the nuclei of cancer cells. Rho-A protein expression was decreased by the effect of 15 μ M curcumin in normal and malignant cell lines (Fig. 5). This effect was more pronounced in malignant and tumorigenic Alpha5 and Tumor2 cell lines. The protein expression of Rho-A in malignant Alpha3 cell line was downregulated in comparison to the control cell line.

Effect of curcumin on DNA damage signaling: PARP-1 determination. DNA damage response is a complex process involving multiple DNA repair cell cycle proteins. To determine the effect of curcumin on DNA damage response PARP-1, an important single strand break binding protein, was analyzed (Fig. 6A). PARP-1 protein is cleaved in response to DNA damage in cells undergoing apoptosis and shows that 30 μ M of curcumin stimulated cleaved PARP-1 protein expression in control MCF-10F, estrogen, Alpha3 and Tumor2. In the case of Alpha5 cell line there was a decrease in cleavage by curcumin treatment. Collectively, these results suggest that curcumin induces apoptosis in breast cancer cells.



Figure 2. (A) Effect of curcumin treatment (0 and 30 μ M) on anchorage-independent growth assay in MCF-10F, Estrogen, Alpha3, Alpha5 and Tumor2 cell lines. Soft agar method was used. Colonies were analyzed after 21 days. (B) Effect of curcumin treatment (0 and 30 μ M) on cell cycle analyses in MCF-10F, Estrogen, Alpha 3, Alpha5 and Tumor2 cell lines. The percentage of the cells entering the specific phase of the cell cycle was determined by flow cytometric analysis of propidium iodide-stained cells.



Figure 3. Effect of curcumin treatment (0 and 15 μ M) on (A) PCNA protein expression by immunoperoxidase method in MCF-10F, Alpha3, Alpha5 and Tumor2 cell lines and (B) on Ras-GRF1 protein expressions by immunofluorescence method in MCF-10F, Alpha5 and Tumor2 cell lines; the graph represents the relative grade of luminescence to assess the protein levels by confocal microscope studies coupled with a computer Adobe Photoshop Program.



Figure 4. Effect of curcumin treatment (0 and 15 μ M) on Ras-GRF1 protein expression determined by Western blot analyses in MCF-10F, Estrogen, Alpha3, Alpha5 and Tumor2 cell lines, β -actin was used as control for loading. Graph represents the relative grade of luminescence to assess the protein level by Western blotting of the cell lines.



Figure 5. Effect of curcumin treatment (0 and 15 μ M) on Rho-A protein expressions determined by Western blot analyses of MCF-10F, Estrogen, Alpha3, Alpha5 and Tumor2 cell lines. β -actin was used as control for loading. The graph represents the relative grade of luminescence to assess the protein level of the cell lines.

Effect of curcumin on DNA damage signaling: γ -H2AX determination. DSBs are generated during apoptosis when chromosomal DNA is cleaved into oligonucleosomal pieces and phosphorylated H2AX (γ -H2AX), forms discrete foci at the sites of DSBs, facilitating the remodeling complexes to the sites of DNA damage, and influences both the efficiency and fidelity of DSB repair. Differential expression levels of γ -H2AX were observed after curcumin treatment (Fig. 6B). It was found in the whole cell lysate, that the γ -H2AX protein expression detected in the MCF-10F, Alpha5 and Tumor2 was diminished upon curcumin treatment. There was no expression of phospho γ -H2AX in the Alpha3 cell line with or without curcumin. However, there was an increase by curcumin in the Estrogen cell line. These results suggest that γ -H2AX foci are stimulated in malignant and transformed Alpha5 and Tumor2 cell lines. On the other hand, the γ -H2AX appear activated by curcumin treatment only in the Estrogen cell line, which possess a normal phenotype.

Discussion

Although accumulating evidence suggests that curcumin has a diverse range of molecular targets (5-8), effects of curcumin in normal and malignant cells are still largely unclear. To understand the effect of curcumin on breast carcinogenesis, we used an *in vitro* breast cancer model system in this study. The present study indicates that curcumin had a significant inhibitory effect on cell growth and colony formation in breast carcinogenesis and is able to activate DNA damage signaling.

Cell cycle analysis of these cell lines showed that S and G₂/M phases were altered by curcumin by reducing the proportion of cells in G_0/G_1 phase and increasing the number of G_2/M phase cells indicating that curcumin imposed a stronger G₂/M checkpoint as compared to untreated cells but less in Tumor2. The effect was more pronounced in Alpha3 and Alpha5 than in Tumor2 cells. That curcumin seems to inhibit cell proliferation, induce apoptosis, and promote accumulation of cells in the G_2/M phase of the cell cycle has been demonstrated by several authors (14-18). Although cell cycle modulators are designed to target cancer cells, some of these can also be applied for a different purpose, as to protect normal cells against the lethality of chemotherapy (14). In some cell types like thymocytes, curcumin induced apoptosis-like changes whereas in many other normal and primary cells curcumin either stimulates or inhibits proliferation. Inhibition of both proliferation and apoptosis of T lymphocytes by curcumin led authors to conclude that the inhibition of cell proliferation by curcumin was not always associated with programmed cell death (14). Interestingly, curcumin has been found to inhibit proliferation of normal, non-selectively, as well as malignant cells, although its apoptogenic effect is more profound in malignant cells since it selectively induced apoptosis in deregulated cyclin D1-expressed cells at G₂ phase of cell cycle in a p53-dependent manner (10,25).

Ras proteins must first release bound GDP mediated by GRF to reach their active GTP bound state. Curcumin increased GRF1 protein expression of MCF-10F and E cell line with or without curcumin preventing the binding. The cell lines Alpha3, Alpha5 and Tumor2 had increased protein expression. However, curcumin decreased such expression increasing the binding.Results also showed that Rho-A protein expression decreased in the malignant cell lines Alpha3, Alpha5 and Tumor2 with 15 μ M curcumin but not in MCF-10F and Estrogen.

Previously, our laboratory demonstrated (24,30-32) that the parental MCF-10F cell line exposed to double doses of



Figure 6. Effect of curcumin treatment (0 and 30 μ M) on (A) cleaved PARP1 and (B) phospho H2AX and H2AX protein expressions in the cell lines: MCF-10F, Estrogen, Alpha3, Alpha5 and Tumor2 cell lines by Western blot analyses. β -actin was used as control for loading.

alpha-particle radiation and treated with estrogen showed a more complex pattern of allelic imbalance compared to cell lines treated with a single dose of radiation without estrogen. Exposure of cell lines to double doses of radiation without estrogen and analyzed at different passages also showed a progressive change. Allelic alterations induced by irradiation with either a single or double dose of alpha particles in the presence or absence of estrogen were expressed either in the form of loss of heterozygosity/microsatellite instability or by some phenotypic changes, such as anchorage independence or invasive capabilities. Consequently, the doses of radiation of the cell lines after irradiation directly influenced these alterations and this genetic effect was more deleterious when given in combination with estrogen. The c-Ha-ras oncogene, mapped to 11p15.5, acquires transforming capacity either by single point mutation (s) in codon 12 or 61, resulting in the expression of an aberrant gene product, or by over-expression of the normal c-Ha-ras p21 protein. Studies in rats have implicated c-Ha-ras mutations in the etiology of breast adenocarcinomas (25,27-29) It is known that estrogen is a key requirement for the normal development of the mammary gland. Although, some reports have shown estrogen to affect c-Ha-ras expression (26).

The present results showed that PARP-1 was cleaved upon curcumin treatment in malignant and tumorigenic cells. Cleaved PARP-1 protein was increased in the presence of curcumin in the control MCF-10F and Estrogen cell lines, as well as in the malignant Alpha3 and the tumor cell line Tumor2. However, it was reduced in Alpha5 cell line indicating that PARP-1 was cleaved upon curcumin treatment in malignant and tumor cells. Curcumin (26) induced apoptosis in tumor cells by activating caspase-8, which leads to activation of caspase-9 and -3, and activation of PARP-1 and apoptosis of tumor cells. Pretreatment with curcumin enhanced the cleavage of PARP-1 in an ovarian cancer cell line after exposure with cisplatin indicating induction of apoptosis (32,33).

The γ -H2AX protein expression was overexpressed in MCF-10F and diminished upon curcumin treatment. There was no expression of γ -H2AX in the Estrogen cell line or Alpha3. However, there was an increase by curcumin effect in the expression in Alpha5 and Tumor2 cell lines by curcumin. Thus, DNA double-strand breaks induced histone H2AX phosphorylation as reported by others (35). Therefore, it can be concluded that PARP-1 and H2AX confer cellular protection against radiation and estrogen-induced DNA damage when curcumin is present.

Targeting either PARP-1 or histone H2AX may provide an effective way of maximizing the therapeutic value of antioxidants for cancer prevention. It is important to point out that curcumin had an effect only on the malignant cell lines and not on the controls, indicating that it can be considered as an important substance for prevention of breast cancer. It can be concluded from these studies that curcumin had significant inhibitory effect on cell growth in breast carcinogenesis. Alteration of multiple protein expressions involved in key signaling pathways render this model an important tool for monitoring the effects of natural dietary compounds in breast carcinogenesis.

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