

Inhibitory effects of fucoxanthinol on the viability of human breast cancer cell lines MCF-7 and MDA-MB-231 are correlated with modulation of the NF-kappaB pathway

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Abstract Fucoxanthin is a carotenoid present in the chloroplasts of brown seaweeds. When ingested, it is metabolized mainly to fucoxanthinol in the gastrointestinal tract by digestive enzymes. These compounds have been shown to have many beneficial health effects. The present study was designed to evaluate the molecular mechanisms of action of fucoxanthin and/or of its metabolite fucoxanthinol against viability of estrogen-sensitive MCF-7 and estrogen-resistant MDA-MB-231 breast cancer cell lines. Fucoxanthin and fucoxanthinol reduced the viability of MCF-7 and MDA-MB-231 cells in dose- and time-dependent manners as a result of increased apoptosis. Furthermore, fucoxanthinol-induced apoptosis was more potent than that of fucoxanthin and correlated, for MDA-MB-231 cells, with inhibitory actions on members of the NF- κ B pathway p65, p50, RelB, and p52. Being overexpressed and regulated by NF- κ B in different types of cancers, the transcription factor SOX9 was also decreased at the nuclear level by fucoxanthin and fucoxanthinol in MDA-MB-231. Taken together, the current results suggest that fucoxanthinol and fucoxanthin could be potentially effective for the treatment and/or prevention of different types of cancers, including breast cancer.

Keywords Fucoxanthin · Fucoxanthinol · MCF-7 · MDA-MB-231 · Viability · Apoptosis · PARP · p65

Introduction

Breast cancer is the most common cancer diagnosed in women worldwide. In the United States alone, it is estimated that 40,000 women will die of breast cancer and over 208,000 new cases will be diagnosed each year (Jemal et al. 2010). These statistics emphasize the urgent need for improvements in detection, diagnosis, and treatment of breast cancer. Several studies have linked diets rich in carotenoids with reduced risk of chronic diseases and cancers, including breast cancer (Lordan et al. 2011; Peng et al. 2011; Tanaka et al. 2012).

Fucoxanthin (Fx) is a naturally occurring brown- or orange-colored pigment that belongs to the class of non-provitamin A carotenoids present in the chloroplasts of brown seaweeds. It is the most abundant of all carotenoids, accounting for more than 10 % of the estimated total natural production of carotenoids (Matsuno 2001). It forms a complex with chlorophyll–protein and plays an important role in light harvesting and photoprotection for effective light use and upregulation of photosynthesis. Oral administration of Fx does not exhibit toxicity and mutagenicity under experimental conditions in mice (Beppu et al. 2009a, b). Carotenoids and Fx, being mostly fat soluble, follow the same intestinal absorption path as dietary fat. Ingested Fx is metabolized mainly to

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fucoxanthinol (Fxl), which is further converted to amarouciaxanthin A in the liver (Asai et al. 2004), thus the bioactive forms of Fx are Fxl and/or amarouciaxanthin A. Dietary Fx is hydrolyzed to Fxl in the gastrointestinal tract by digestive enzymes such as lipase and cholesterol esterase and then absorbed into intestinal cells (Sugawara et al. 2002). Fx and Fxl have been shown to have many beneficial health effects, including anti-mutagenic (Nishino et al. 2009), antidiabetic (Nishikawa et al. 2012), antiobesity (Maeda et al. 2005), anti-inflammatory (Kim et al. 2010b; Shiratori et al. 2005), and preventive actions on liver, breast, prostate, colon, and lung cancers (Das et al. 2005; Kotake-Nara et al. 2005; Le Marchand et al. 1993; Nishino et al. 2009; Slattery et al. 2000; Zhang et al. 1999). Most of these actions involve modulation of the nuclear factor kappaB (NF- κ B) signaling pathway.

The transcription factor members of the NF- κ B family found in most cells are p50, p65/RelA, c-Rel, p52, and RelB (Hayden and Ghosh 2004). NF- κ B is regulated by two main pathways: the canonical and noncanonical pathways (Hayden and Ghosh 2008). In its resting state, NF- κ B dimers are secured in the cytoplasm by I κ B proteins which render the NF- κ B transcription factors inactive. These I κ B members α , β , and ϵ contain two conserved serine residues that become phosphorylated by I κ B kinases (IKKs) upon NF- κ B pathway activation, followed by proteasome-dependent degradation and NF- κ B transcription factors activation (Karin and Ben-Neriah 2000). NF- κ B members then translocate to the nucleus to form dimers likely to have distinct regulatory functions. Among the NF- κ B target genes are inflammatory cytokines, anti-apoptotic proteins, and cell cycle regulators that promote cancer cells proliferation (Karin and Lin 2002). Indeed, NF- κ B is activated by most carcinogens, leading to expression of anti-apoptotic gene products which allow survival and growth of tumors (Baker et al. 2011; Kumar et al. 2004; Zubair and Frieri 2013). It has been shown that p65 and p50 subunits of NF- κ B are constitutively active and are overexpressed in breast cancer (Nakshatri et al. 1997), resulting in further transcription of anti-apoptotic genes (Fan et al. 2008).

The objective of the current research was to determine if Fx and/or Fxl have inhibitory effects on viability and on the NF- κ B signaling pathway of breast cancer cell lines. Comparative studies have been undertaken in two different types of cells, the MCF-7 and MDA-MB-231 cell lines. Since MCF-7 cells inherently express

high levels of estrogen receptor (ER)- α and MDA-MB-231 express very low, if any, ER- α or - β , we sought to determine if the presence of a functional ER altered the effects of Fx and/or Fxl treatments. Here, we report that Fxl has been shown to reduce cell viability, to increase apoptosis, and to regulate components of the NF- κ B pathway in both cell lines from breast cancers.

Material and methods

Chemicals

Fucoxanthin (Fx) and fucoxanthinol (Fxl) were purchased from Wako Chemicals (Richmond, VA, USA).

Cell culture

Human breast cancer cell lines MCF-7 (hormone sensitive) and MDA-MB-231 (hormone resistant) were obtained from the American Type Culture Collection (ATCC; Bethesda, MD, USA). MCF-7 and MDA-MB-231 cells were grown in DMEM/F12 medium supplemented with 10 % heat-inactivated fetal bovine serum (Canadian origin) and 100 U/ml penicillin/streptomycin (Corning, Tewksbury, MA, USA). Cells were cultured at 37 °C and 5 % CO₂.

Cell viability

The viability of Fx- or Fxl-treated cells were measured using CellTiter-Blue cell viability assay (Promega, Madison, WI, USA) according to manufacturer's instructions. Following dose (20, 30, or 40 μ M) and time-dependent (12, 24, or 48 h) treatments with Fx or Fxl of breast cancer cell lines, cell viability was estimated by measuring the amount of reduced resorufin by its fluorescence at 560_{Ex}/590_{Em} using a multimode microplate reader (Varioskan, Thermo Scientific, Waltham, MA, USA). IC₅₀ values were determined from linear regression of fluorescence versus concentration and correspond to doses of Fx or Fxl necessary to reduce cell viability by 50 % for 12 h of treatment.

Apoptotic, necrotic, and healthy cells assay

The breast cancer cell lines were treated with 20 μ M Fx or Fxl for 12 h, followed by staining with FITC-Annexin V, ethidium homodimer III, and Hoechst

33342 according to the manufacturer's protocol (Biotium, Inc., Hayward, CA, USA). Fluorescence was assessed using an Axio Observer A1 inverted fluorescence microscope (Carl Zeiss, Gottingen, Germany) with FITC, rhodamine, and DAPI filters. Images were merged and analyzed using the ImageJ software (<http://rsbweb.nih.gov/ij/>).

Western blot analysis

The breast cancer cell lines were treated with 20 μM Fx or Fxol for 12 h, followed by protein extractions for nuclear and cytoplasmic proteins, using hypertonic buffer with phosphatase inhibitors (10 mM NaF, 1 mM Na_3VO_4 , and 20 mM glycerol2-phosphate) and protease inhibitors (30 $\mu\text{g}/\text{ml}$ aprotinin, 2.5 $\mu\text{g}/\text{ml}$ leupeptin, 10 $\mu\text{g}/\text{ml}$ pepstatin, and 1 mM phenylmethyl fluoride) as described previously (Schreiber et al. 1989). Protein concentrations of protein extracts were evaluated using the Bradford method (Bradford 1976). Ten micrograms (for nuclear extracts) or 30 μg (for cytoplasmic extracts) of protein from each sample was separated by SDS–polyacrylamide gels (SDS–PAGE) and electrophoretically transferred onto a PVDF membrane. Membranes were incubated in blocking solution containing 5 % nonfat dry milk in TBST buffer (TBS buffer containing 0.1 % Tween 20) for 1 h at room temperature, followed by incubation with a specific primary antibody overnight at 4 °C. Specific proteins were detected using the following primary antibodies: polyclonal anti-CASPASE 3 (1:1,000, Cat.: 9662, Cell Signaling, Danvers, MA, USA), polyclonal anti-Poly(ADP-ribose) polymerase-1 (PARP) (1:1,000, Cat.: 9542, Cell Signaling), NF- κB family member antibody sampler kit (1:1,000, Cat.: 4766, Cell Signaling), monoclonal anti-Phospho-NF- κB p65 (Ser536) (1:1,000, Cat.: 3033, Cell Signaling), monoclonal anti-STAT3 (1:2,000, Cat.: 4904, Cell Signaling), polyclonal anti-STAT5 (1:1,000, Cat.: 9363, Cell Signaling), polyclonal anti-SOX9 (1:500, Cat.: AB5535, Millipore, Billerica, MA, USA), monoclonal anti- α -Tubulin (1:5,000, Cat.: 05-829, Millipore), and polyclonal anti-NCL (1:1,000, Cat.: 12247S, Cell Signaling). The membranes were washed three times in TBST buffer for 5 min, followed by incubation for 1 h with horseradish peroxidase-conjugated secondary antibody. The membranes were washed again and developed using an enhanced chemiluminescent detection system (Luminata Forte, Millipore) according to the manufacturer's instructions. Images were taken using the FluorChem IS 8900 imaging system (Alpha Innotech,

San Jose, CA, USA). The Western blotting results were referred to NCL (for nuclear extracts) and α -Tubulin level (for cytoplasmic extracts) as references.

Statistics

Experiments for cells viabilities were repeated at least three times, and the data were presented as means \pm S.E.M. Statistical analysis of the data was performed using Student's *t* test and ANOVA with GraphPad Prism (GraphPad Software Inc., San Diego, Ca, USA). $P < 0.05$ was considered significant.

Results

Fx and Fxol inhibit cell viability in human breast cancer cells

We first examined the effects of Fx and Fxol on cell viability of human breast cancer cell lines MCF-7 and MDA-MB-231 (Fig. 1). Fx and Fxol significantly decreased cell viability in a time- and dose-dependent manner for both cell lines. A two-way ANOVA test showed a significant effect of time and dose on cell viability ($P < 0.0001$). Fxol-induced suppression of cell viability was more rapid and more pronounced, compared to Fx. Indeed, IC_{50} values of 121.89 and 39.63 μM for Fx and Fxol, respectively, were calculated for MCF-7, whereas for MDA-MB-231, IC_{50} values of 141.54 and 33.59 μM for Fx and Fxol were determined for 12 h of treatment. In addition, these results also demonstrate a more efficient inhibitory action of Fxol on cell viability of MDA-MB-231, compared to MCF-7, for all times of treatments investigated. Only Fxol had significant inhibitory effects on viability at 12 h with doses of 30 and 40 μM of treatments in both breast cancer cell lines. Although giving nonsignificant viability reduction, we used the dose of 20 μM for 12 h as treatments with Fx or Fxol for the remaining of the study to obtain enough cells for protein extractions and the study of apoptotic markers by Western blot assays.

Reduction of breast cancer cell lines viability in response to Fxol involves increase of apoptotic pathways activities (Fig. 2). In MCF-7 cells, lacking caspase-3 expression (Jänicke 2009), PARP was cleaved in response to 20 μM Fx and Fxol stimulations for 12 h and resulted in cytoplasmic (Fig. 2a) and nuclear (Fig. 2b) accumulations of the cleaved fragment.

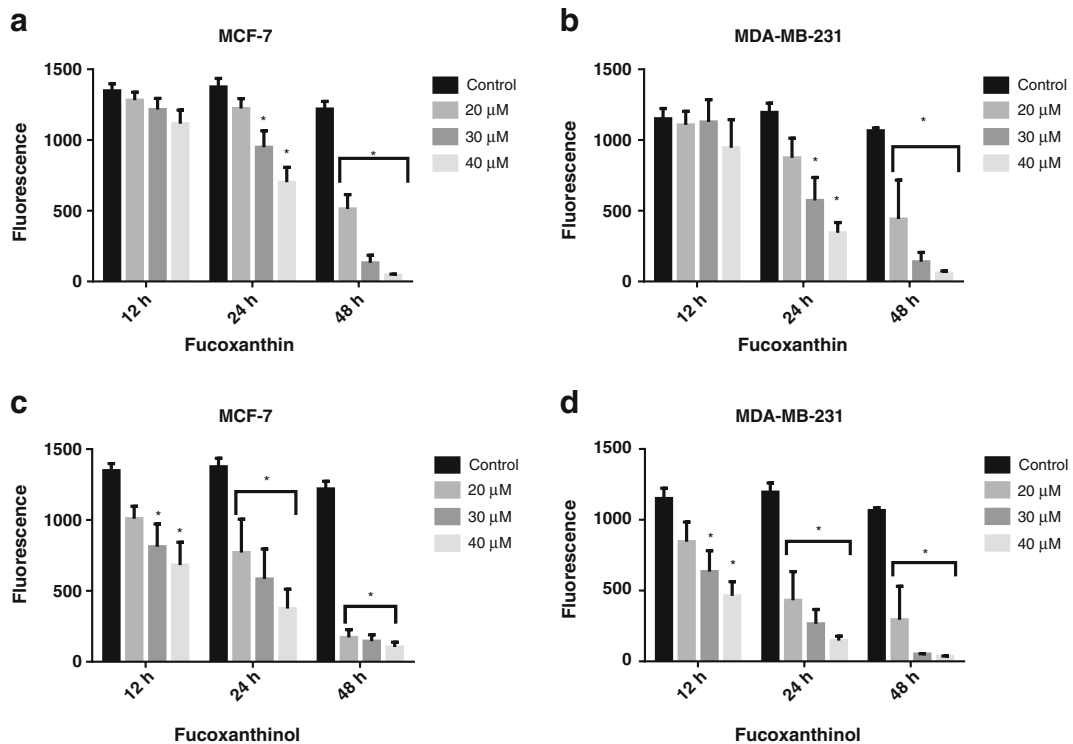


Fig. 1 Fx and Fxol reduce cell viability of MCF-7 and MDA-MB-231 cell lines. Exponentially growing cells were incubated in the absence or presence of 20–40 μM of Fx or Fxol, for 12–48 h, followed by determination of cell viability as described in [Material and methods](#). The results are presented as absolute fluorescence

signal and are expressed as means \pm standard error of mean deviation of four independent experiments in which each treatment was performed in quadruplicate. Comparisons to control were done using a multiple comparison test (*significant, $P < 0.05$)

For MDA-MB-231 cells, procaspase-3 was cleaved following Fxol treatment (Fig. 2a), whereas PARP cleavage in response to Fx and Fxol treatments resulted in nuclear accumulation of the cleaved form (Fig. 2b) with no cytoplasmic detection. Increased apoptosis of breast cancer cell lines in response to treatments for 12 h with 20 μM Fx or Fxol was also confirmed using fluorescence microscopy (Fig. 2b). For both cell lines, Fxol resulted in higher apoptotic levels compared to Fx treatment as shown with FITC-Annexin V green fluorescence (Fig. 2c). However, although a major difference was observed between Fx and Fxol on viability reduction of MDA-MB-231 cells (Fig. 1b, d), Fxol treatment resulted in a slightly higher level of apoptosis when compared to Fx (Fig. 2c, lower panels). In both cell lines, Fx and Fxol treatments did not result in increased levels of necrosis as shown with ethidium homodimer III red fluorescence. Thus, these changes correspond to increased apoptosis of breast cancer cells in response to Fx/Fxol.

Regulatory effects of Fx and Fxol on the NF- κB signaling pathway

Since several NF- κB target genes are involved in inhibiting apoptosis in breast cancer cells (Nakshatri and Goulet 2002; Wu and Kral 2005; Zhou et al. 2005a), we examined whether Fx and/or Fxol inhibits members of the NF- κB pathway. In MCF-7 cells, Fxol, but not Fx, increased the nuclear levels of p65, p50, cRel, and p52 (Fig. 3b). In addition, the nuclear levels of phosphorylated p65 increased in response to Fxol. Since cytoplasmic concentrations of p65, p50, and p52 remained constant following Fxol treatments of MCF-7 cells (Fig. 3a), increased nuclear accumulations of these NF- κB members might be attributed to increases in their protein syntheses. However, decreased cytoplasmic concentrations of cRel in response to Fxol in MCF-7 cells suggest that its nuclear accumulation is mainly attributed to increased nuclear translocation. In MDA-MB-231 cells, Fxol contributed to decreased nuclear

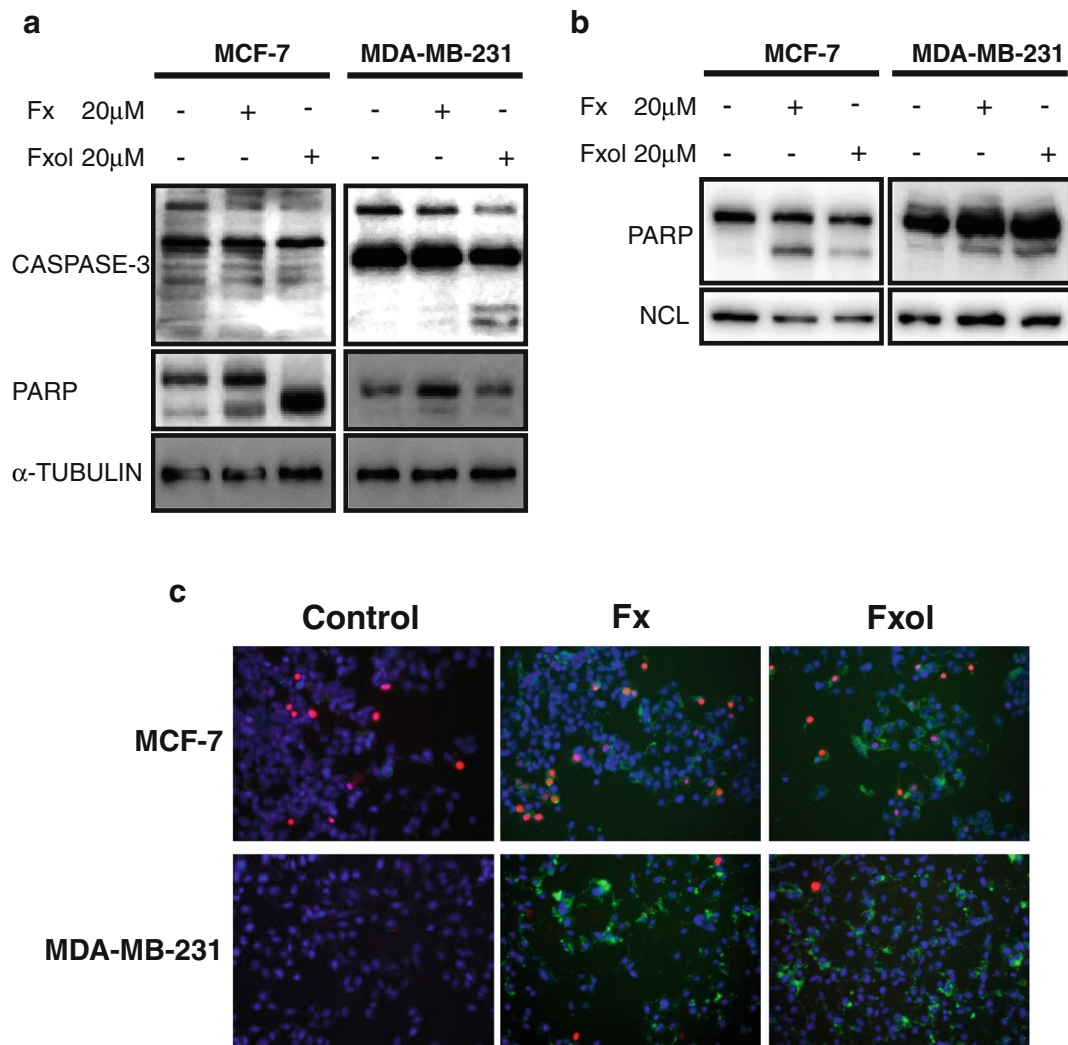


Fig. 2 Effects of Fx and Fxol on markers of apoptosis in breast cancer cell lines. MCF-7 and MDA-MB-231 cell lines were treated with 20 μ M Fx or Fxol for 12 h. Cytoplasmic (a) and nucleic (b) proteins were extracted from control and treated cells, followed by determination of Caspase 3 and PARP cleavages using Western blot analysis as described in [Material and methods](#) and use of α -Tubulin and Nucleolin (NCL) as loading controls for cytoplasmic

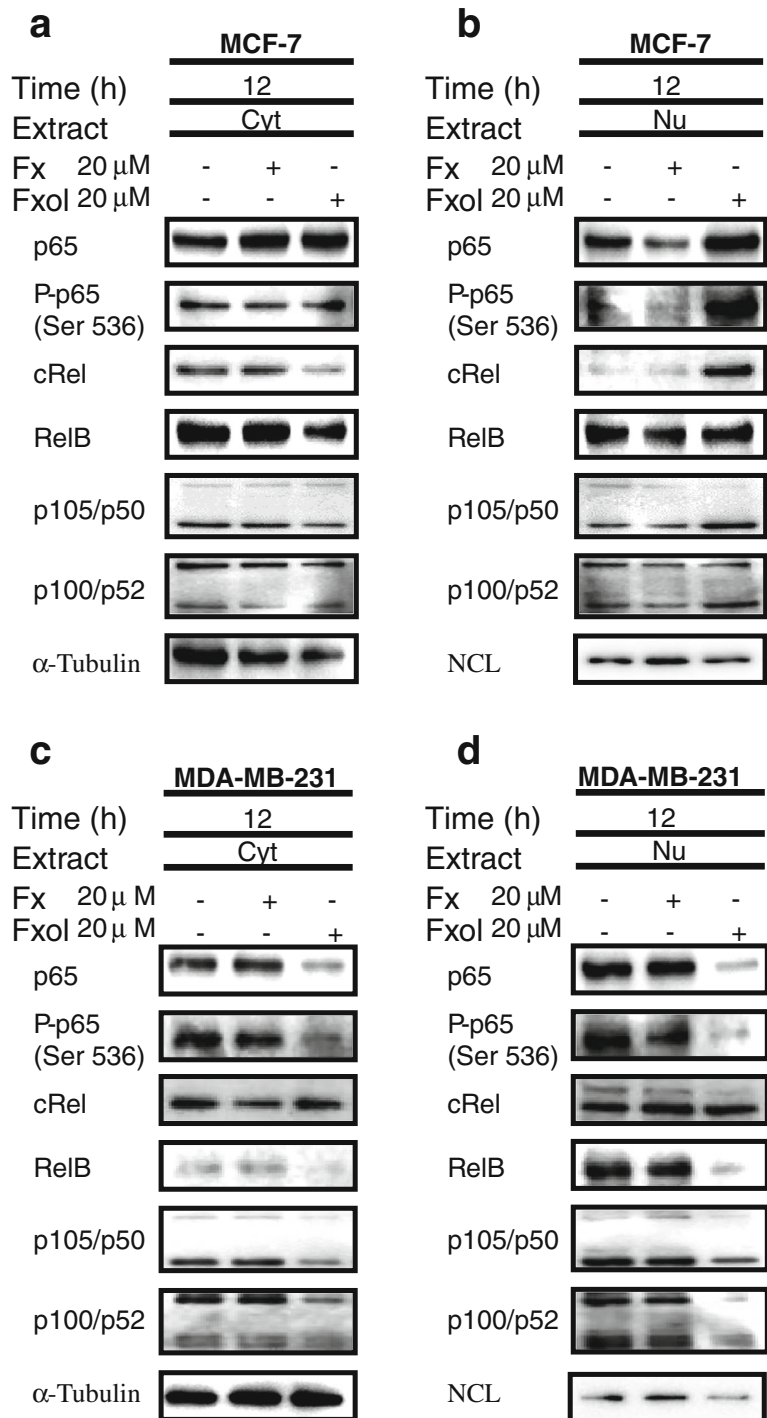
and nuclear extracts, respectively. In c, MCF-7 cells (*left panel*) and MDA-MB-231 cells (*right panel*) were stained with Hoechst 33342 (for viable cells in *blue*), FITC-Annexin V (for apoptotic cells in *green*), and ethidium homodimer III (for necrotic cells in *red*), followed by imaging with an inverted fluorescence microscope with total magnifications of $\times 200$. The experiments were repeated three times

levels p65, p50, RelB, p52/p100, and phosphorylated-p65 (Fig. 3d). In this cell line, decreased nuclear levels of components of the NF- κ B pathways were attributed to decreased synthesis, as shown from cytoplasmic protein extracts (Fig. 3c).

Since SOX9 has been shown to be regulated by NF- κ B at the transcriptional level in other cancer cells (Saegusa et al. 2012; Sun et al. 2013), we investigated whether Fx and/or Fxol could influence SOX9 expression in breast cancer cell lines. Although no effects on SOX9 could be

observed in MCF-7 cells (Fig. 4a), SOX9 expression and nuclear accumulation were decreased in response to Fx and Fxol in MDA-MB-231 cells (Fig. 4b). SOX9 Western blots from MDA-MB-231 extracts show the presence of two forms of the protein. Such result might be attributed to SOX9 phosphorylation, as previously shown by protein kinase A in Sertoli-like cells (Malki et al. 2005). Therefore, Fx and Fxol treatments might result in loss of SOX9 phosphorylation, as seen with the disappearance of the upper band (Fig. 4b).

Fig. 3 Effects of Fx and Fxol on members of the NF- κ B pathway. MCF-7 and MDA-MB-231 cell lines were treated with 20 μ M Fx or Fxol for 12 h and cytoplasmic and nuclear proteins were extracted from controls and treated cells. The levels of target proteins were determined using Western blot analysis as described in [Material and methods](#). α -Tubulin and Nucleolin (*NCL*) were used as loading controls for cytoplasmic and nuclear extracts, respectively. Every experiment was repeated three times



Influences of Fx and Fxol on members of the STAT family

Since deregulation of STAT signaling cascades is associated with cellular proliferation and transformation

(Kisseleva et al. 2002), we investigated whether Fx and/or Fxol could have any effects on expression and nuclear localization of members of the STAT family in breast cancer cells. Although Fxol had an inhibitory effect on STAT5 protein synthesis in MCF-7 cells

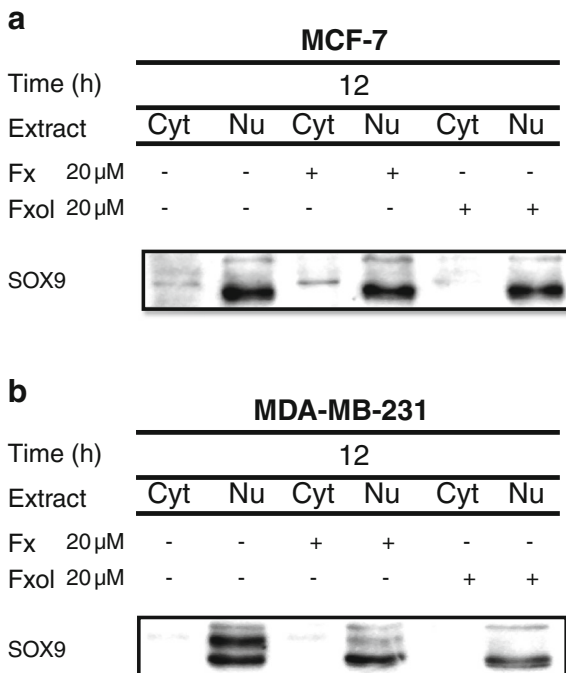


Fig. 4 Effects of Fx and Fxol on SOX9 protein levels and nuclear translocation. MCF-7 and MDA-MB-231 cell lines were treated with 20 μ M Fx or Fxol for 12 h and cytoplasmic and nuclear proteins were extracted from controls and treated cells. The levels of SOX9 were determined using Western blot analysis as described in [Material and methods](#). Every experiment was repeated three times

(Fig. 5a), no changes in nuclear concentrations for STAT3 and STAT5 in response to Fx or Fxol could be observed in MCF-7 and STAT5 did not vary in MDA-MB-231 nuclear extracts (Fig. 5b, d). Unlike others (Lahusen et al. 2007; Yuan et al. 2010), we were not able to detect STAT5 at the protein level in MDA-MB-231 cells. However, Fx alone had an inhibitory effect on cytoplasmic accumulation of STAT3 in MDA-MB-231 (Fig. 5c), suggesting that nuclear translocation of STAT3 is increased in response to Fx in MDA-MB-231 cells to maintain a constant nuclear concentration (Fig. 5d). Together, these results suggest that inhibitory effects of Fx and Fxol on cell viability may be independent of the STAT signaling pathway.

Discussion

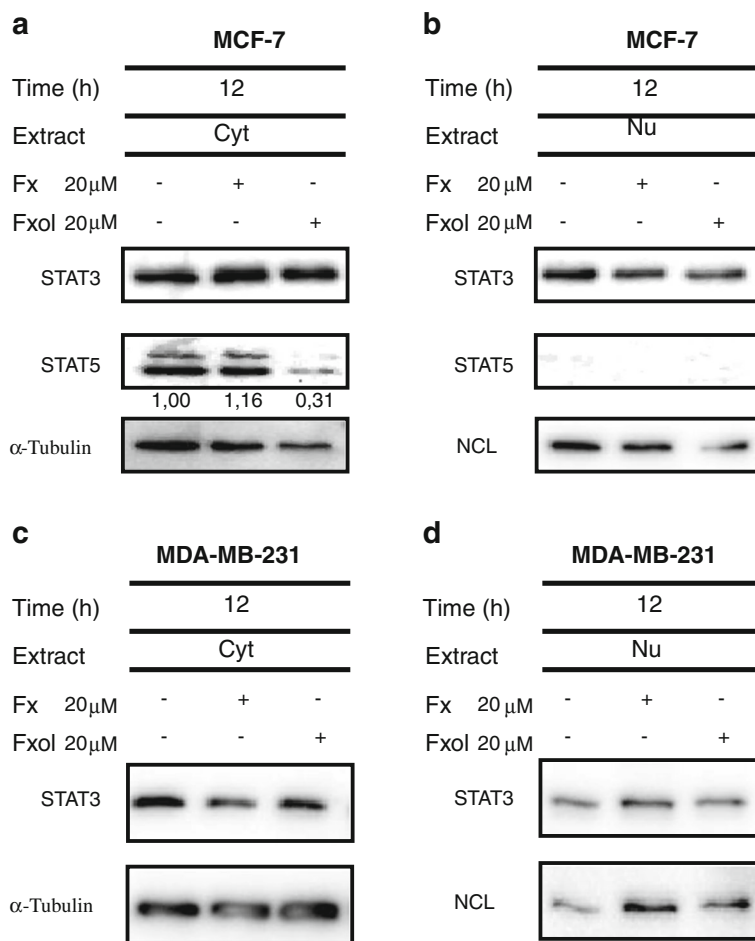
In the current study, we demonstrated that Fxol show inhibitory effects on cell viability and apoptosis-

inducing effects on two human breast cancer cell lines, MCF-7 and MDA-MB-231. Furthermore, the apoptosis-inducing activities of Fxol were more potent than that of Fx and were correlated, for hormone independent MDA-MB-231 cells, to inhibitory actions on members of the NF- κ B pathway.

As shown in primary effusion lymphoma cell lines (Yamamoto et al. 2011), Fxol-induced suppression of cell viability was more pronounced than that of Fx for breast cancer cell lines MCF-7 and MDA-MB-231. Compared to Fx, Fxol also had a stronger antiproliferative effect on PC-3 human prostate cancer cells (Asai et al. 2004), MCF-7 and Caco-2 human colon cancer cells (Konishi et al. 2006), thus supporting our current results on breast cancer cells. Indeed, Fxol has been shown previously to reduce MCF-7 cells viability by 75 % following treatment with 25 μ M Fxol for 48 h (Konishi et al. 2006). Differences with our results (87 % reduction with 20 μ M Fxol for 48 h) may be attributed to variations in culture conditions and Fxol concentration, purity, and stability.

Hallmarks of the apoptotic process are cleavages of PARP, an enzyme implicated in DNA damage and repair mechanisms, and of Caspase-3, an enzyme activated during apoptosis both by extrinsic (death ligand) and intrinsic (mitochondrial) pathways. During apoptosis, PARP is cleaved from its precursor having a mass of 116 kDa to yield an 85 kDa fragment (Kaufmann et al. 1993). PARP fragment migrates from the nucleus into the cytoplasm during late stages of apoptosis associated with severe nuclear fragmentation (Soldani et al. 2001), suggesting that apoptosis is more advanced in MCF-7 compared to MDA-MB-231 cells treated with Fxol. Others have shown that Fx induced Caspase-3, -7, and PARP cleavages and thus triggered apoptosis of human leukemia HL-60 and prostate cancer PC-3 cells (Kim et al. 2010a; Kotake-Nara et al. 2005). Fx and Fxol also induced cell cycle arrest in G1 phase and caspase-dependent apoptosis, inhibited the activation of NF- κ B, and downregulated anti-apoptotic proteins in primary effusion lymphoma cells (Yamamoto et al. 2011). The present study clearly demonstrated that Fxol induced apoptosis of breast cancer cells MDA-MB-231 and MCF-7 by cleavages of procaspase-3 and PARP. In support to our results, Fxol isolated from *Halocynthia roretzi* induced apoptosis in human breast cancer cells (Konishi et al. 2006). Lack of difference in necrosis levels between treatments in breast cancer cells suggests that Fx and Fxol have no effects on necrotic

Fig. 5 Effects of Fx and Fxol on STAT members' protein levels and nuclear translocation. MCF-7 and MDA-MB-231 cell lines were treated with 20 μ M Fx or Fxol for 12 h and cytoplasmic and nuclear proteins were extracted from controls and treated cells. The levels of STAT3 and STAT5 proteins were determined using Western blot analysis as described in [Material and methods](#). Results for STAT5 in MCF-7 were analyzed for densitometry using the ImageJ software (<http://rsbweb.nih.gov/ij/>) and expressed as a ratio between STAT5 and Tubulin followed by normalization to control. Every experiment was repeated three times



pathways. Indeed, apoptosis and necrosis are two processes than can occur independently, sequentially, or simultaneously (Elmore 2007; Zeiss 2003). Presence of necrosis in MCF-7 cells may be attributed to suboptimal cell culture conditions. Thus, our results suggest that reduction of breast cancer cells viability in response to Fx and Fxol treatments may be attributed to increased apoptosis.

Estrogen receptors play an important role in breast cancer; women with ER-positive tumors have an overall better prognosis and are more likely to have their tumors respond to therapy. However, in up to 25 % of cases, ER-positive tumors are nonresponsive to therapy as a result of acquired resistance, possibly linked to constitutive NF- κ B leading to estrogen-independent growth (Fraser et al. 2009; Kalaitzidis and Gilmore 2005; Zubair and Frieri 2013). In this study, we looked at possible differences in the inhibitory actions of Fx and/

or Fxol on components of the NF- κ B pathway between estrogen-sensitive MCF-7 and estrogen-resistant MDA-MB-231 breast cancer cell lines. In lipopolysaccharide (LPS)-stimulated RAW 264.7 macrophages, others have shown that Fx inhibited the nuclear translocation of p50 and p65 proteins, resulting in lower levels of nuclear transactivation by NF- κ B transcription factors (Kim et al. 2010b; Lee et al. 2013). In the present study, major differences were observed in the inhibitory mechanisms between Fx and Fxol on the activation of members of the NF- κ B pathway in breast cancer cell lines. Indeed, Fxol treatment decreased nuclear levels p65, p50, RelB, and p52/p100 in hormone independent MDA-MB-231 cells. Thus, Fxol treatment may contribute to reduce viability of aggressive estrogen-independent tumor growth and may involve inhibitions of nuclear translocation and transcriptional activity of members of the NF- κ B pathway. Indeed, other NF- κ B inhibitors such

as celastrol and triptolide have been shown to reduce MDA-MB-231 cell viability (Kang et al. 2009; Kannaiyan et al. 2011; Liu et al. 2009). Further regulation of NF- κ B members by phosphorylation, acetylation, and coactivator and corepressor interactions in response to Fx/Fxol remains to be investigated. Surprisingly, our results show increased nuclear localization of members of the NF- κ B pathway (p65, p50, cRel) in response to Fxol in MCF-7 as opposed to MDA-MB-231 cells. Although others have shown that inhibition of NF- κ B activation in MCF-7 cells results in reduced cell viability (Zhou et al. 2005b), our results suggest that decreased MCF-7 cells viability in response to Fx/Fxol may be independent of NF- κ B inhibition. However, the balance between the expression of genes involved in the survival and apoptotic processes ultimately determines whether the cell undergoes apoptosis in response to NF- κ B (Aggarwal 2000). Indeed, treatment of MCF-7 cells with Fxol resulted in increased apoptosis despite the fact that NF- κ B transcription factors were activated, thus suggesting that other signaling pathways may be involved.

In breast cancer cells, SOX9, being regulated by NF- κ B in other cell types (Saegusa et al. 2012; Sun et al. 2013), has been reported mainly in the cytoplasm of MDA-MB-231 cells, and was located mostly in the nucleus of MCF-7 cells (Chakravarty et al. 2011). We showed that nuclear levels of SOX9 were unchanged in MCF-7 cells when stimulated with Fx or Fxol, whereas SOX9 was located mainly in the nucleus of MDA-MB-231 cells. These differences with the Chakravarty et al. 2011 study may be attributed to differences in cell culture and potential cross-reaction of the antibody they used against SOX9. Interestingly, upregulation of SOX9 in response to retinoic acid has also been shown to induce cell growth arrest in breast cancer cells (Müller et al. 2010). This is in contrast to our results where treatments with Fx and Fxol resulted in decreased nuclear levels of SOX9. Indeed, SOX9 may rather be involved in MDA-MB-231 cells proliferation and downregulation of its phosphorylation may contribute to the inhibitory effects of Fx and Fxol on viability of estrogen resistant breast cancers.

In addition to having crucial roles in mammary gland development, STAT3/5 transcription factors also contribute to tumorigenesis when deregulated (Desrivières et al. 2006; Kisseleva et al. 2002). Indeed, constitutive activation of STAT3 has been linked to the growth and survival of human breast cancer cells (Garcia et al. 1997,

2001). Higher concentrations of Fx (50–150 μ M) than what was used in the current study have been shown to inhibit STAT3 expression and phosphorylation in sarcoma tissue (Wang et al. 2012) and gastric adenocarcinoma cells (Yu et al. 2011). Although we have shown that Fx and Fxol had no effect on nuclear levels of STAT3/5 in breast cancer cells, the phosphorylation profiles of these transcription factors may be modulated to promote apoptosis. Indeed, decreased phosphorylation of STAT3 is associated to reduced invasiveness and survival of breast tumors (Hsieh et al. 2005).

In conclusion, Fx and Fxol effectively induced cell viability reduction and apoptotic cell death of breast cancer cells. Thus, dietary Fx or Fxol could be potentially effective for the treatment and/or prevention of different types of cancers, including breast cancer. Indeed, most dietary Fx may be converted to Fxol, and Fxol may exert a suppressive effect on cancer cells more efficiently than Fx in vivo. However, although there is a correlation between increased apoptosis and decreased nuclear accumulations of p65, RelB, p50, and p52 in MDA-MB-231 cells, more research will be required to clearly establish a regulatory action of Fxol on members of NF- κ B to activate apoptosis.

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Conflict of interest The authors declare that there is no conflict of interest that would prejudice their impartiality.

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