

Arachidonic and eicosapentaenoic acids induce oxidative stress to suppress proliferation of human glioma cells

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Introduction

The most malignant tumor of the brain is glioblastoma multiforme (commonly called gliomas). Despite debulking surgery, radiation and chemotherapy, the survival of patients diagnosed to have glioblastoma multiforme is not more than 44 weeks. Gliomas are difficult to treat, partly because while growing they merge with normal brain tissue and hence during surgery it is difficult to delineate the tumor tissue from normal brain tissue to excise them completely. In view of this, developing newer therapeutic strategies that target glioma cells selectively with minimal toxicity to normal brain cells is urgently needed. Previous studies performed by us and others revealed that polyunsaturated fatty acids (PUFAs), especially γ -linolenic acid (GLA, 18:3 n-6), arachidonic acid (AA, 20:4 n-6), eicosapentaenoic acid (EPA, 20:5 n-3) and docosahexaenoic acid (DHA, 22:6 n-3), have selective tumoricidal action and induce apoptosis of glioma cells both *in vitro* and *in vivo* [1–7].

Previous studies revealed that many tumor cells have decreased expression of Δ^6 and Δ^5 desaturases that are essential for the formation of long-chain metabolites of dietary essential fatty acids (EFAs): linoleic (LA, 18:2 n-6) and α -linolenic (ALA, 18:3 n-3) acids. This decrease in the activity of desaturases results in a deficiency of long-chain metabolites of EFAs such as GLA, dihomo-GLA (DGLA, 20:3 n-6), AA formed from LA and EPA and DHA from ALA in the tumor cells, possibly to protect themselves (tumor cells) from the cytotoxic action of PUFAs, free radicals (generated during the metabolism of EFAs/PUFAs) and lipid peroxides derived from various PUFAs [reviewed in 8, 9]. In a previous preliminary open label clinical study, we showed that intratumoral infusion/injection of GLA regresses glioblastoma [3, 4, 6] suggesting that some PUFAs can be exploited for the therapy of cancer including gliomas. This assumption is supported by the studies performed in cell cultures, rodent glioma and other tumor models, and preliminary human studies [1–19].

The tumoricidal action of various PUFAs has been attributed to their ability to enhance free radical generation and lipid peroxides specifically in tumor cells, changes in the lipid content of the cell membrane due to the incorporation of supplemented PUFA, action on anti-angiogenic factors and enzymes involved in lipid metabolism, changes in P-gly-

coprotein expression and induction of changes in mitochondrial function [1–19]. It is known that the expression of different oncogenes differently affects sensitivity and/or resistance of various cancer cells to the cytotoxic action of anti-cancer drugs [17, 18]. In a previous study [20], we showed that AA produces its tumoricidal action on IMR-32 (human neuroblastoma) cells by enhancing the expression of *FAS* and caspases 3 and 8 compared to the control, indicating the involvement of the extrinsic apoptotic pathway [21]. However, it is not known whether the same mechanism plays a role in the induction of death of other human-derived neuroblastoma cells in the presence of PUFAs. Hence, we performed the present study to explore this possibility by studying the effect of various PUFAs on two different human glioma cells (LN229, HNGC2) *in vitro*.

The results of this study revealed that AA and EPA suppress *p53*, *Ras*, *Myc*, *BCL-2/Bax*, *NF-κB/κB* and *5-LOX* and enhance cytochrome C and caspases 3 and 9 expression secondary to an increase in the accumulation of lipid peroxides and thus inhibit tumor cell growth.

Material and methods

Chemical reagents

Cell culture compounds including DMEM high glucose culture medium (Cat. No: 12100-046) and heat inactivated fetal bovine serum from Gibco (Cat. No: 16000-044) were obtained from Incell technologies Pvt. Ltd, India; penicillin – streptomycin (Cat. No: P0781), amphotericin (Cat. No: A2942), trypsin EDTA (Cat. No: T4049), MTT (Cat. No: M5655), COX (Cat. No: I7378) and LOX (Cat. No: 74540) inhibitors, RTq-PCR (Cat. No: 1318855) components were from Sigma Aldrich Pvt. Ltd (Bangalore, India); primers from Bioserve Hyderabad, India. All fatty acids were procured from Cayman Chemical Company (California, USA).

Cell culture conditions

Glioblastoma cell lines LN229 and HNGC2 were obtained from the Centre for Cellular and Molecular Biology (CCMB) Hyderabad, India. Both the cell lines were subcultured in DMEM (pH 7.4) medium containing NaHCO₃, 100 U/ml penicillin, 100 µg/ml streptomycin, 1.25 µg/ml amphotericin B and 10% fetal bovine serum (FBS). Cells seeded in vented flasks (T25 cm²) in which they grew as a monolayer in humidified 5% CO₂ ambience at 37°C. On reaching 80% confluence, the cells were washed with phosphate buffered saline (PBS, pH 7.4) and trypsinized (trypsin – 0.25%, EDTA – 0.02%) for 2–3 min. Trypsin was inactivated using FBS. The cells obtained by centrifugation

were tested for their viability using the trypan blue dye exclusion method by counting the cells in a hemocytometer. The pellets of cells obtained were passaged and used for further experimental studies [19, 20].

Cell proliferation assay

Effect of various fatty acids on glioma cells

Glioma cells LN229 and HNGC2 were seeded at a density of 5×10^3 cells/100 µl/well in 96-well culture plates. The cells were allowed to attach for 24 h and supplemented with fresh medium and used for further studies.

Effect of n-6 and n-3 PUFAs on growth of glioma cells

Glioma cells LN229 and HNGC2 were seeded at a density of 5×10^3 cells/100 µl/well in 96-well culture plates and allowed to attach for 24 h. At the end of the attachment period, cells were treated with different doses of (10, 20, 30 µg/ml) n-6 and n-3 PUFAs and incubated for 24 h. Cell viability was evaluated at the end of the incubation period by MTT assay as described previously [22–24].

Effect of COX and LOX inhibitors on AA/EPA-induced changes in proliferation of glioma cells

Since both AA and EPA form precursors to various prostaglandins, thromboxanes, leukotrienes, lipoxin A4 (from AA) and resolvins (from EPA), we next studied whether COX and LOX inhibitors can influence the actions of AA and EPA on the proliferation of glioma cells LN229 and HNGC2. For this purpose the cells were seeded at a density of 5×10^3 cells/100 µl/well in 96-well culture plates and allowed to attach for 24 h at the end of which they were treated with different doses of COX and LOX inhibitors (indomethacin and NDGA respectively) in the presence of AA/EPA [19, 20].

Effect of AA/EPA on lipid peroxides and nitric oxide

Glioma cells were seeded in 24-well plates at a density of 1×10^5 cells per ml of culture medium per well and allowed to attach for 24 h before the treatment. After the attachment period, cells were treated with AA/EPA (30 µg/ml) as representative of n-6 and n-3 fatty acids respectively for 24 h. After the treatment period, spent medium was collected and cells were washed with 500 µl of PBS (pH 7.4). Washed cells were lysed with NET lysis buffer. Lipid peroxides and nitric oxide levels were measured in both the cell lysates and supernatants as described previously [19, 20].

Effect of AA/EPA on antioxidant enzymes

For this study, LN229, HNGC2 cells were seeded in a 24-well plate at a density of 1×10^5 cells per ml of culture medium per well. The cells were allowed to attach for 24 h before the treatment. After the attachment period, cells were treated for 24 h with AA/EPA (10, 20 and 30 $\mu\text{g/ml}$). At the end of the treatment period, spent medium (supernatant) was collected and used for various estimations. Cells were washed using 500 μl of PBS (pH 7.4) and were lysed using NET lysis buffer. Antioxidant enzymes – superoxide dismutase, catalase, glutathione-s-transferase and glutathione peroxidase – were measured in both the supernatants and cell lysates [19, 20].

Previously, we performed dose- and time-dependent studies with n-6 (LA, GLA, DGLA and AA) and n-3 (ALA, EPA and DHA) on the proliferation of LN229 and HNGC2 cell *in vitro* using MTT assay. In this study, the two glioma cell lines were exposed to 10–50 $\mu\text{g/ml}$ (10, 20, 30, 40, 50 $\mu\text{g/ml}$) of PUFAs for 24 and 48 h. The results of this study showed that all PUFAs have the ability to decrease the proliferation of LN229 and HNGC2 cells *in vitro*. Furthermore, it was observed that a 30 $\mu\text{g/ml}$ dose of both n-6 and n-3 PUFAs at the end of 24 h of the incubation produced an ~50% decrease in proliferation ($p < 0.05$) compared to the control (see Figures 1 A and B for results obtained with 10, 20 and 30 $\mu\text{g/ml}$ of PUFAs tested). In view of these results, in the present study a 30 $\mu\text{g/ml}$ dose of AA and EPA and incubation for a 24 h period were employed. Furthermore, we used in all subsequent studies only AA and EPA as representative of n-6 and n-3 fatty acids.

Molecular gene expression studies

RNA isolation

Glioma cells were seeded at a density of 2×10^6 /ml/well in 6-well culture plates and were used for RNA extraction. After the initial attachment period, glioma cells were treated with AA/EPA (30 $\mu\text{g/ml}$) for 24 h. At the end of the incubation period, TRIzol reagent method was employed to isolate RNA. Quantification and purity of the RNA sample were tested by spectrophotometer (Biochrome libra S70, J.L. Tech, Hyderabad, India). 1 μg of RNA/sample/well was loaded on 1.5% agarose gel to check the integrity of the sample.

C-DNA synthesis (semi-quantitative RT-PCR)

To study various gene expression levels, C-DNA synthesis was performed using SuperScript First-Strand for qRT-PCR. C-DNA was amplified using specific primers by PCR (Eppendorf thermocycler-5331; USA). Human β -actin was chosen as the active con-

trol. The expression of various genes was measured by qRT-PCR as described previously (ref. BBRC). On the completion of PCR, products of each gene processed were separated by means of an electric field at 100 V and size on 1.5% agarose gels in 1x TAE buffer, stained with ethidium bromide. The following primer pairs from 5'-3' were used respectively: *β -actin* – 243 bps: F:CGTGGGCCGCCCTAGGCACCA, R: TTGGCCTTAGGGTTCAGGGGGG. *Bax* – 367 bps: F: ACCAAGAAGCTGAGCGAGTGTC, R: ACAAGATGGTCACGGTCTGCC. *Bcl2* – 319 bps: F: CGACGACTTCTCCCGCCGCTACCGC, R: CCGCATGCTGGGGCCGTACAGTTCC. *P53* – 293 bps: F: CAGCCAAGTCTGTGACTTGACGTAC, R: CTATGTCGAAAAGTGTTCCTGTCATC. *Cyt C* – 242 bps: F: GCGTGTCTTGGACTTAGAG, R: GGCGGCTGTGTAAGAGTATC. *Casp9* – 325 bps: F: GGTCTGGAGATTTGGTGA, R: GACAGCCGTGAGAGAGAATGA. *Casp3* – 378 bps: F: AGAAGATCACGCAAAGGAGC, R: TCAAGCTTGTCCGACTACTG. *NF-kB* – 300 bps: F: AGCACAGATACCACCAAGACCC, R: CCCACGCTGCTCTTCTATAGGAAC. *IKB* – 68 bps: F: TTGGGTGCTGATGTCAATGC, R: AGTCCACTGCGAGGTGAAG. *Ras* – 164 bps: F: TGGAGTCGTAGGACCCTGA, R: TGTGTTCTGATAAGGCCCC. *Myc* – 19 5bps: F: CGACCACAAGGCCCTCAGTA, R: CAGCCTTGGTGTGGAGGAG. *Fos* – 236 bps: F: GAATAAGATGGCTGCAGCCAAATGCCGCAA, R: CAGTCAGATCAAGGGAAGCCACAGACATCT. *COX-I* – 494bps: F: GTTCAACACCTCCATGTTGGTGGAC, R: TGGTGTGAGGCAGACCAGCTTC. *COX-II* – 558bps: F: GTCTGATGATGTATGCCA-CATCTG, R: ATGCCAGTGATAGGGGTGTTAAA. *5-LOX* – 327 bps: F: ATCAGGACGTTACGGCCGA, R: GTCCACGATCTGCTCAATGGT.

PCR cycling conditions: denaturation at 94°C for 30 s, annealing at 52.5°, 63°, 67°, 57°, 62°, 62°, 61°, 62°, 68°, 57°, 58°, 74°, 63°, 64°, 57° for 30 s respectively to the above-mentioned gene sequences and extension at 72° for 2 min, 36 cycles were carried out for amplification. Observed bands for specific genes on the gel were photographed by a gel documentation system which has an UV light trans-illuminator hood and a camera for image capturing. All the genes were quantified using image analysis software (Gel documentation unit, Major science, J.L. Tech, Hyderabad, India).

Statistical analysis

All the above experiments were repeated twice on two different occasions. The data values

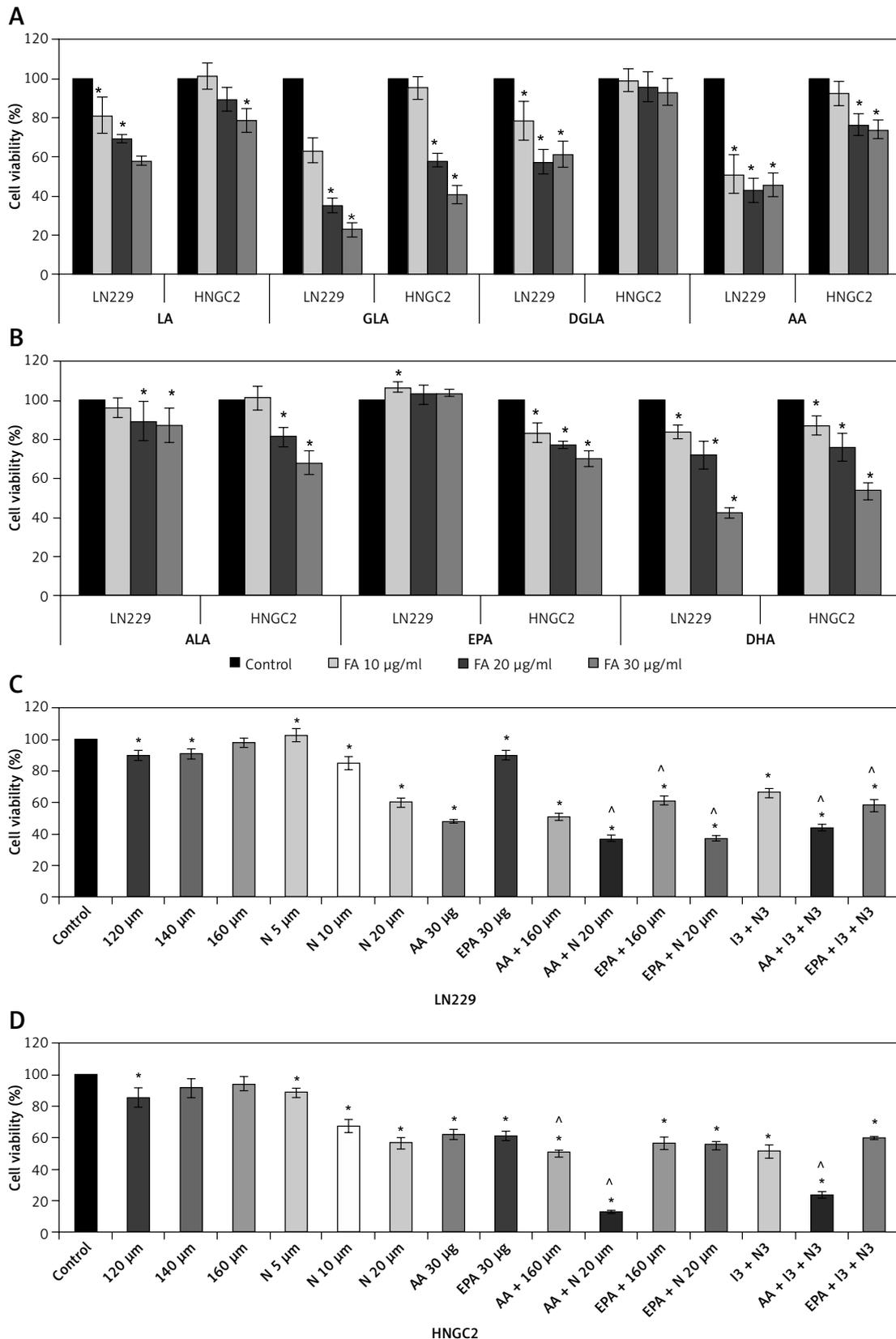


Figure 1. Effect of n-6, n-3 PUFAs on LN229/HNGC2 cells: LN229/HNGC2 cells ($1 \times 10^4/100 \mu\text{l}$) were exposed to different doses of n-6 PUFAs, n-3 PUFAs, and incubated for 24 h. At the end of the treatment period, cell viability was measured by MTT assay. **A** – LN229/HNGC2 cells treated with n-6 PUFAs (10, 20, 30 $\mu\text{g/ml}$). **B** – LN229/HNGC2 cells treated with n-3 PUFAs (10, 20, 30 $\mu\text{g/ml}$). All values are expressed as mean \pm SEM, $n = 6$. * $P < 0.001$, compared to control; **C** – Effect of COX and LOX inhibitors indomethacin and NDGA respectively on growth of LN229/HNGC2 cells. $^{\wedge}P < 0.05$ – compared to respective fatty acid treated (AA/EPA) groups. **D** – Effect of COX/LOX inhibitors \pm AA/EPA on growth of LN229/HNGC2 cells *in vitro*

C – control, FA – fatty acids, AA – arachidonic acid, LA – linoleic acid, GLA – γ -linolenic acid, DGLA – dihomo gamma linolenic acid, EPA – eicosapentaenoic acid, DHA – docosahexaenoic acid, ALA – α -linolenic acid.

showed reproducible results when observed and are expressed as mean \pm SEM. To evaluate the significance of differences between the mean values by paired *T*-test, data were statistically analyzed using the MS Excel statistical analysis tool.

Results

Cell viability assay

Effect of n-6 and n-3 PUFAs on proliferation of glioma cells

When LN229 glioma cells were treated with 10, 20, 30 $\mu\text{g/ml}$ of ω -6 (LA, GLA, DGLA and AA) and ω -3 PUFAs (ALA, EPA and DHA), all PUFAs produced a significant dose-dependent reduction in cell proliferation. Among n-6 PUFAs, LA/GLA/AA (30 $\mu\text{g/ml}$) and among n-3, EPA/DHA (30 $\mu\text{g/ml}$) showed maximum growth inhibitory action. On the other hand, GLA (30 $\mu\text{g/ml}$) and DHA (30 $\mu\text{g/ml}$) exerted maximum growth inhibitory action on HNGC2 cells (Figures 1 B and C). Of all the fatty acids tested, GLA is the most effective growth inhibitory fatty acid on the glioma cells studied. However, we selected AA and EPA for further studies since they form precursors to various eicosanoids and are also able to produce a significant reduction in the proliferation of both the glioma cells studied. It may be mentioned here that PUFAs induced apoptosis of the glioma cells tested (data not shown).

Effect of COX and LOX inhibitors on proliferation of glioma cells

We next studied the effect of the COX inhibitor indomethacin and the LOX inhibitor NDGA on the growth of glioma cells *in vitro*. These studies revealed, in both the glioma cells studied, that the COX inhibitor indomethacin (20, 40, 60 $\mu\text{g/ml}$) by itself did not have any action on the proliferation while the LOX inhibitor NDGA suppressed the proliferation of LN229 and HNGC2 cells *in vitro*. In contrast, both indomethacin and NDGA did not have any effect on the growth inhibitory action of AA and EPA on LN229 and HNGC2 cells. In fact, surprisingly, when the glioma cells were supplemented with indomethacin + NDGA + AA/EPA enhanced suppression of growth inhibitory action of AA/EPA was noted (Figures 1 C and D). These results imply that COX metabolites may not have much significant action on the growth of glioma cells studied. To verify this possibility, we next studied the effect of various prostaglandins and LTD4 and LTE4 (10, 50 and 100 ng/ml) on the growth of LN229 and HNGC2 cells *in vitro*. These results showed that all the prostaglandins tested (PGE1, PGE2, PGI2, PGF2) and leukotrienes (LTD4 and LTE4) suppressed the growth of LN229 and HNGC2 by not

more than 10–25%, which barely showed significance compared to the control (data not shown). These results suggest that, possibly, eicosanoids have insignificant action on the growth of LN229 and HNGC2 cells *in vitro*. Based on these results, we suggest that the growth inhibitory actions of AA and EPA on LN229 and HNGC2 cells noted may be due to the fatty acids themselves with little or no involvement of eicosanoids.

Biochemical assays

Effect of AA/EPA on formation of lipid peroxides and nitric oxide

The results of these studies showed that both nitric oxide and lipid peroxides were increased in the supernatants and cell lysates of glioma cells in the presence of AA/EPA thus can augment oxidative stress in glioma cells (LN229 > HNGC2, Table I). It is evident that the total levels of lipid peroxides (LPO) (supernatant + cell lysate) were significantly elevated in LN229 cells in the presence of both AA and EPA. However, in the HNGC2 cells AA treatment did not produce any significant increase in the production of lipid peroxides (both in the supernatant and cell lysate) compared to the control. On the other hand, EPA did induce a significant increase in lipid peroxides in both the supernatant and cell lysate compared to the control.

Effect of AA/EPA on antioxidant enzymes in glioma cells

The results of the studies performed with AA/EPA revealed that both AA and EPA produced a significant increase in the activity of catalase, SOD, glutathione-s-transferase and glutathione peroxidase when compared to the control. These results suggest that AA/EPA produce significant changes in the antioxidant defenses of glioma cells (LN229 > HNGC2) (Table I B).

Gene expression studies (semi-quantitative RT-PCR)

Effect of AA (30 $\mu\text{g/ml}$)/ EPA (30 $\mu\text{g/ml}$) on mRNA expression of various genes

It is known that the apoptotic signaling pathway consists of an extrinsic (death receptor pathway) and an intrinsic component (mitochondrial pathway) [25]. *Bcl-2* family proteins play a vital role in the survival of cells by acting on the caspases of the intrinsic pathway. In view of this, we studied the effect of AA/EPA on the expression of *BCL-2* and caspases and other factors that influence cell proliferation and apoptosis.

These results revealed that both the glioma cells treated with AA and EPA produced a significant

Table I. Effect of AA/EPA (30 µg/ml) on nitric oxide and lipid peroxides production and antioxidant enzymes in LN229 and HNGC2 cells. Both the glioma cells were treated with AA/EPA (30 µg/ml) and incubated for 24 h. At the end of treatment, spent medium was collected and cells were lysed. **A** – Both cell lysate and spent medium were used for measurement of nitric oxide and lipid peroxides. **B** – Cell lysate was used for measurement of anti-oxidant enzymes

A

Parameter	LPO [µM]			NO [µM]		
	Supernatant	Lysate	Lysate + supernatant	Supernatant	Lysate	Lysate + supernatant
LN229:						
Control	1.102 ±0.012	0.298 ±0.001	1.400 ±0.011	1.110 ±0.021	0.760 ±0.020	1.866 ±0.002
AA 30 µg	1.427 ±0.035*	0.271 ±0.006	1.699 ±0.031*	1.410 ±0.083	0.700 ±0.033	2.116 ±0.100*
EPA 30 µg	1.195 ±0.034*	0.404 ±0.002	1.598 ±0.035*	1.290 ±0.078	0.766 ±0.076	2.066 ±0.073*
HNGC2:						
Control	1.102 ±0.061	0.282 ±0.006	1.384 ±0.006	1.216 ±0.031	1.000 ±0.173	2.216 ±0.082
AA 30 µg	1.058 ±0.034	0.249 ±0.007*	1.308 ±0.038	1.050 ±0.037*	0.717 ±0.049	1.767 ±0.055*
EPA 30 µg	1.355 ±0.041*	0.406 ±0.002*	1.762 ±0.043	1.116 ±0.048	0.816 ±0.028	1.933 ±0.031*

B

Parameter	Catalase [µM H ₂ O ₂ /min/gm protein]	SOD [U/mg protein]	GST [µM/min/gm protein]	GPX [µM/min/gm protein]
LN229:				
Control	9009.36 ±1358.9	184.011 ±43.03	18.144 ±3.325	24503.38 ±2727.16
AA 30 µg	16404.15 ±2289*	906.59 ±211*	68.80 ±16.42*	78451.58 ±22350.09
EPA 30 µg	9830.72 ±2870.25	686.12 ±211.02	38.02 ±6.67*	23481.96 ±4937.79
HNGC2:				
Control	9821.87 ±2287	127.44 ±44.5	18.23 ±4.94	35106.08 ±1042.45
AA 30 µg	11565.24 ±2616	393.75 ±85	26.49 ±3.95	60053.54 ±5906.215*
EPA 30 µg	1000.17 ±1693.52	245.56 ±60.58	19.60 ±0.79	46316.38 ±4699.59

All values are expressed as mean ± SEM. *P < 0.05 – compared to control, AA – arachidonic acid, EPA – eicosapentaenoic acid.

decrease in *BCL-2/Bax* ratio (Figure 2 A). In a similar fashion, AA/EPA produced a significant decrease in the expression of *NF-κB/κB* (Figure 2 B), implying that AA/EPA suppress inflammation. In contrast to these results, AA/EPA enhanced the expression of cytochrome C and caspases 3 and 9 significantly in both the glioma cells (AA > EPA) (Figure 2 C) compared to the control.

AA and EPA induced a significant decrease in the expression of *p53*, *Ras* and *Myc* (except that AA and EPA did not have any significant effect on *Ras* expression in HNGC2 cells, Figures 3 A and B) in both the glioma cells compared to the control. In contrast, a significant increase in expression of *Fos* was seen in both LN229 and HNGC2 cells when exposed to AA and EPA (Figures 3 A and B), though HNGC2 cells when supplemented with EPA 30 µg/ml dose showed significantly decreased expression of *Fos*.

Neither of the glioma cells showed any change in *COX-1* expression on exposure to AA and EPA. Ironically, *COX-2* expression was significantly enhanced by AA and EPA in both the glioma cells. In contrast, the expression of *5-LOX* was also signifi-

cantly decreased in both the glioma cells by AA and EPA in comparison to the control (Figure 3 C).

To understand and compare the effects of AA and EPA on various indices studied in the present study on both the glioma cells, and their summary is depicted in Table II.

Discussion

Our results presented here suggest that both LN229 and HNGC2 glioma cells are sensitive to the growth inhibitory action of LA/GLA/AA and EPA/DHA (Figures 1 A and B). These results are similar to those we obtained with IMR-32 cells [20]. LN229 cells showed greater sensitivity to the growth inhibitory action of PUFAs compared to HNGC2 cells. Despite the fact that the LOX inhibitor NDGA, but not indomethacin, a COX inhibitor, suppressed the proliferation of LN229 and HNGC2 cells *in vitro*, various prostaglandins and LTs tested showed only marginal growth suppressive action on LN229 and HNGC2 cells (data not shown). These results indicate that the growth inhibitory action of AA and EPA on LN229 and HNGC2 cells

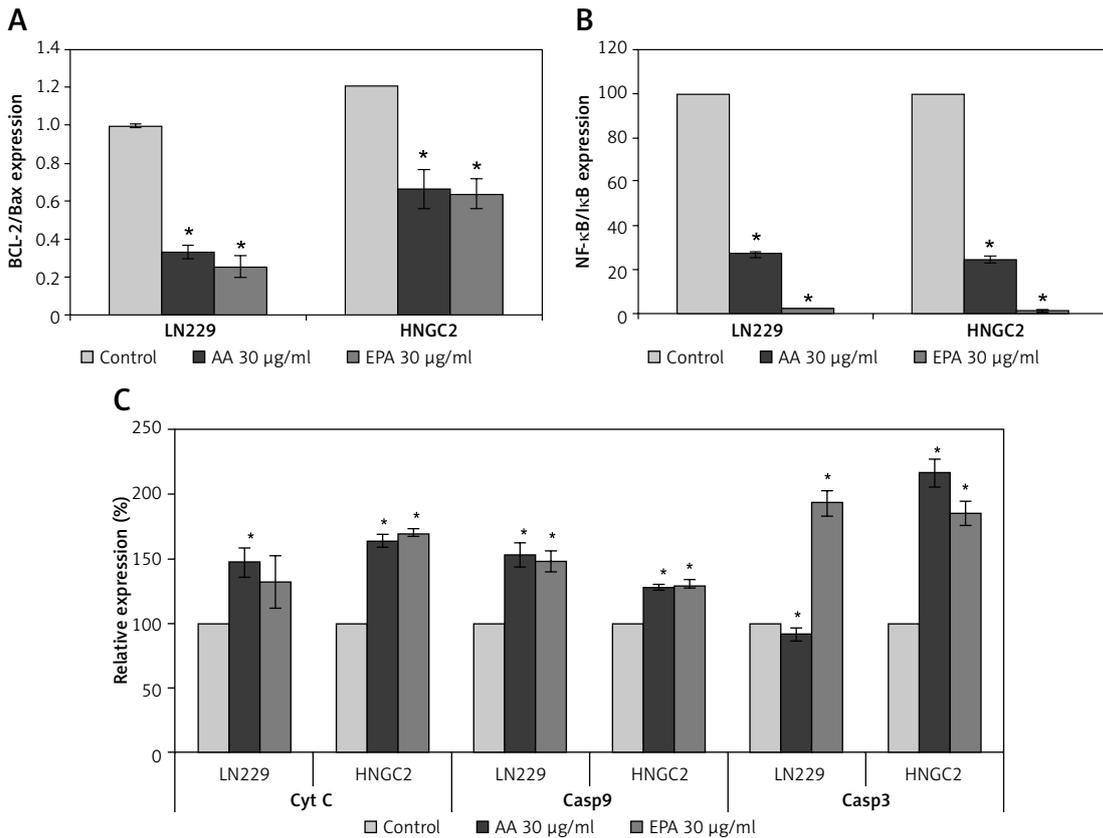


Figure 2. Effect of AA/EPA (30 µg/ml) on mRNA gene expression of Bcl-2-Bax, inflammatory genes NF-κB/IκB and apoptotic genes cyt C, casp9, casp3 in LN229 and HNGC2 cells. **A** – Expression of Bcl-2/Bax ratio in LN229 and HNGC2 cells. **B** – Expression of NF-κB/IκB ratio in LN229 and HNGC2 cells. **C** – Expression of cyt C, caspase 9 and caspase 3 genes in LN229 and HNGC2 cells. The beta actin gene was loaded as a positive control

All values are expressed as mean ± SEM. *P < 0.05 compared to control; AA – arachidonic acid, EPA – eicosapentaenoic acid, CytC – cytochrome C, Casp9 – caspase 9, Casp3 – caspase 3.

may be due to the fatty acids themselves with little or no role for eicosanoids.

Further evaluation of the possible mechanism(s) of action of AA and EPA on glioma cells revealed that the *Bcl-2/Bax* ratio (Bax is essential for apoptosis whereas overexpression of *Bcl-2* enhances cell survival) was tilted more towards *Bax* (AA > EPA; LN229 > HNGC2), suggesting an increase in the rate of apoptosis (Figure 2 A). The *Bcl-2/Bax* ratio is much lower in EPA treated cells compared to AA treatment, which may explain why AA is more effective than EPA in inhibiting the growth of LN229 compared to HNGC2 cells. Both AA and EPA were almost equally effective in suppressing the *NF-κB/IκB* ratio in both LN229 and HNGC2 cells (Figure 2 B). In contrast, a significant increase in the expression of *Fos* was seen in both LN229 and HNGC2 cells when exposed to AA and EPA (Figures 3 A and B), though HNGC2 cells when supplemented with EPA in a 30 µg/ml dose showed significantly decreased expression of *Fos*.

Neither of the glioma cells showed any change in *COX-1* expression but showed significantly enhanced expression of *COX-2* and *5-LOX* by AA and EPA. These results could be interpreted to mean

that glioma cells may produce more eicosanoids that do not have any significant action on their proliferation and thus reduce the intracellular concentration of toxic AA and EPA to avoid apoptosis. Furthermore, increased production of eicosanoids may augment glioma cell proliferation since inflammation has pro-carcinogenic action [22–24].

Overall, the most significant observation of the present study is an increase in the formation of lipid peroxides and NO and increased expression of cytochrome C and caspases 3 and 8 (Figure 2 C, Table I) by AA and EPA in both the glioma cells examined. These results suggest that AA and EPA enhance the formation of lipid peroxides in tumor cells, which may result in an increase in the expression of cytochrome C and caspases 3 and 8, leading to their apoptosis. Thus, AA and EPA induce apoptosis of glioma cells by activating both the extrinsic and intrinsic apoptotic pathways. A close examination of the results obtained in the present study suggests that the more the lipid peroxides generated in the tumor cells, the higher the concentrations of antioxidants and higher the rate of apoptosis (Figures 1 A, B, 2 C and Table I),

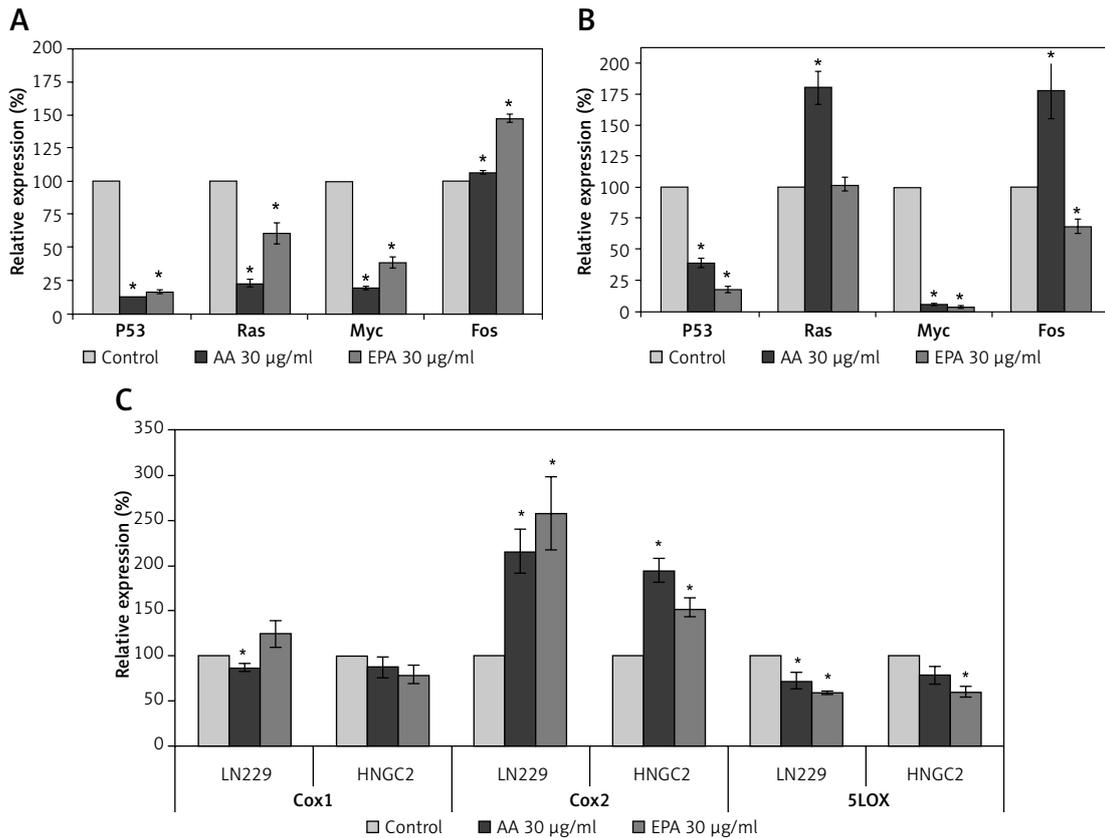


Figure 3. Effect of AA/EPA (30 µg/ml) on mRNA gene expression of oncogenes and metabolic genes COX-1, COX-2, 5-LOX in LN229 and HNGC2 cells. **A** – Expression of P53, Ras, Myc Fos, in LN229 cells. **B** – Expression of P53, Ras, Myc Fos, in HNGC2 cells. **C** – Expression of Cox1, Cox2 and 5LOX in LN229 and HNGC2 cells. The β-actin gene was loaded as a positive control. All values are expressed as mean ± SEM. **P* < 0.05 compared to control. It may be noted here that there are no error bars for the controls in Figures 1–3 since we calculated the cell viability for the control group as 100%. In these figures error bars for the control groups are not visible because we compared and calculated the control group values with the control group itself, so the error is the value ‘0’. Also, the treated group values were compared and calculated with the control group values, resulting in error bars
C – control, *AA* – arachidonic acid, *EPA* – eicosapentaenoic acid

which suggests why AA is more potent than EPA in inducing apoptosis of LN229 cells compared to HNGC2 cells *in vitro* and LN229 cells show greater sensitivity to the growth inhibitory actions of AA compared to HNGC2 cells. This may also explain the slightly higher expression of caspase 9 seen in LN229 cells compared to HNGC2 cells (Table II, though cytochrome C and caspase 3 expression levels are almost the same in both the glioma cells). Though we have not specifically investigated whether there is any direct correlation between lipid peroxides generated, degree of apoptosis and sensitivity of glioma cells to PUFAs, this interesting possibility needs to be evaluated in future studies. We aware of the limitations of the present study since we did not perform measurements of real-time PCR of p53, Ras, Myc, BCL-2/Bax, NF-κB/ IκB, and 5-LOX, FACS results of annexin and pi staining and Western blot of caspases to establish apoptosis of glioma cells on exposure to PUFAs. Despite this limitation, previously, we and others have shown [1, 2, 8–13, 16] that PUFAs indeed

produce apoptosis of tumor cells on incubation with various PUFAs at the doses used in the present study, observations that support our present results and conclusions reached.

It is known that apoptosis is induced by the extrinsic death receptor pathway and the intrinsic mitochondrial pathway. Activation of caspases leads to the onset of apoptosis. Notably, both pathways converge at caspase-3, leading to activation of other proteases. The results of our study showed that in both the glioma cell lines, AA and EPA increased cytochrome C and caspase-3 and caspase-9 but decreased the *Bcl-2/Bax* ratio. Under normal physiological conditions, expression of *Bax* is increased in response to tumor suppressor protein *p53*, and *Bax* is involved in *p53*-mediated apoptosis. The *p53* protein is a transcription factor, which is a part of the cell’s response to stress and regulates many downstream target genes, including *Bax*. Apoptosis is regulated in part by the *Bcl-2* gene, which promotes cell survival. *Bax* overexpression enhances apoptosis by

Table II. Summary of effects of AA/EPA on various indices studied in glioma cells *in vitro*

Fatty acid	Parameter	LN229	HNGC2
AA	Cell proliferation	↓↓↓	↓↓
EPA		↔	↓↓
AA	Bcl-2/Bax	↓↓↓	↓↓
EPA		↓↓	↓
AA	NF-κB/IκB	↓↓	↓↓
EPA		↓↓↓	↓↓↓
AA	Cytochrome C	↑↑	↑↑
EPA		↑↑	↑↑
AA	Caspases 3 and 9	↑↑	↑↑
EPA		↑↑	↑↑
AA	P53	↓↓↓	↓↓
EPA		↓↓	↓↓
AA	Ras	↓↓↓	↑↑
EPA		↓	↔
AA	MYC	↓↓↓	↓↓↓
EPA		↓↓	↓↓↓
AA	FOS	↑	↑↑
EPA		↑↑	↓
AA	COX-1	↓	↔
EPA		↔	↔
AA	COX-2	↑↑	↑↑
EPA		↑↑↑	↑↑
AA	5-LOX	↓	↔
EPA		↓↓	↓↓

the intrinsic pathway. Thus, the balance between *Bcl-2* and *Bax* genes is an important determinant of apoptosis. In the present study, it is observed that *p53* expression and the *Bcl-2/Bax* ratio were decreased (in other words, *Bcl-2* is decreased and *Bax* is increased. i.e. relatively there is decreased expression of *Bcl-2* and relatively higher expression of *Bax*). It is known that *p53* interacts with *Bax*, decreasing its activation to induce apoptosis. The results of the present study showed that both the glioma cell lines, when treated with AA and EPA, showed decreased *p53* and *Bax* expression levels, which is contrary to the regulatory effect of *p53* on *Bax*. Generally, it is expected that *p53* has positive control of *Bax* expression, i.e. if *p53* expression is increased it leads to an increase in *Bax* expression as well. In contrast, in the present study, we noted the reverse, namely, a decrease in

p53 expression but an increase in *Bax* expression. This suggests uncoupling of the relationship between *p53* and *Bax* expression.

Taking all the results of the present study into consideration, it is evident that AA and EPA suppressed *p53*, *Ras*, *Myc*, *BCL-2/Bax*, *NF-κB/IκB* and *5-LOX* and enhanced cytochrome C and caspases 3 and 9 expression and increased the accumulation of lipid peroxides and nitric oxide and altered their antioxidant defenses (LN229 > HNGC2). These results suggest that possibly enhancement in the formation of lipid peroxides and consequent oxidative stress is responsible for the growth inhibitory action of AA and EPA and other PUFAs on glioma cells. These results are supported by our previous studies wherein we observed that AA induced apoptosis of IMR-32 cells, a human neuroblastoma cell line that is similar in characteristics to HNGC2 and LN229, by enhancing lipid peroxides and oxidative stress [20]. In the previous study [20] we noted that IMR-32 cells undergo apoptosis as a result of activation of the extrinsic apoptotic pathway due to the expression of *Fas* and caspases 8 and 3. In the present study, we did not study the expression of *Fas* and caspase 8 and so the involvement of the extrinsic pathway in the apoptosis of LN229 and HNGC2 cells remains conjectural.

In conclusion, it is known that caspases play a significant role in inflammation, cell proliferation, tumor suppression, cell differentiation, neural development and axon guidance and ageing [25] that may explain pleiotropic actions of AA and EPA and their metabolites [26, 27].

Conflict of interest

The authors declare no conflict of interest.

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