

Lipids, Mitochondria and Cell Death: Implications in Neuro-oncology

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Received: 5 April 2010 / Accepted: 5 April 2010 / Published online: 29 April 2010
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Abstract Polyunsaturated fatty acids (PUFAs) are known to inhibit cell proliferation of many tumour types both in vitro and in vivo. Their capacity to interfere with cell proliferation has been linked to their induction of reactive oxygen species (ROS) production in tumour tissues leading to cell death through apoptosis. However, the exact mechanisms of action of PUFAs are far from clear, particularly in brain tumours. The loss of bound hexokinase from the mitochondrial voltage-dependent anion channel has been directly related to loss of protection from apoptosis, and PUFAs can induce this loss of bound hexokinase in tumour cells. Tumour cells overexpressing Akt activity, including gliomas, are sensitised to ROS damage by the Akt protein and may be good targets for chemotherapeutic agents, which produce ROS, such as PUFAs. Cardiolipin peroxidation may be an initial event in the release of cytochrome *c* from the mitochondria, and enriching cardiolipin with PUFA acyl chains may lead to increased peroxidation and therefore an increase in apoptosis. A better understanding of the metabolism of fatty acids and eicosanoids in primary brain tumours such as gliomas and their influence on energy balance will be fundamental to the possible targeting of mitochondria in tumour treatment.

Keyword Lipids · Mitochondria · Cell death · Neuro-oncology

Introduction

Gliomas are the most common form of central nervous system tumour, and the high-grade, malignant gliomas (World Health Organization grades III and IV) are notoriously difficult to treat. They exhibit a highly aggressive phenotype with invasive behaviour, which permits rapid spread throughout the brain, although they rarely form metastases [1]. Long-term survival rates for malignant glioma patients have improved very little with the advent of impressive improvements in both brain tumour diagnosis and surgery. Thus, despite many treatment options including surgical debulking, radiotherapy, and more recently chemotherapy, patient prognosis remains dismal and advances in novel treatment strategies are as urgently needed.

Polyunsaturated fatty acids (PUFAs) are known to inhibit cell proliferation of many different tumour types both in vitro and in vivo. Their capacity to interfere with cell proliferation has been linked to their induction of reactive oxygen species (ROS) production in tumour tissues leading to cell death, but the exact mechanisms of action are far from clear, particularly in brain tumours including gliomas. This review provides a brief summary of the effects of PUFAs upon mitochondrial metabolism, cell proliferation and apoptosis and raises questions about the mechanisms by which PUFAs exert their antitumour effects on gliomas in experimental models.

Fatty Acid Effects on Tumour Cell Proliferation

PUFAs have been found to inhibit the cell proliferation of many different tumour types in vitro. The earliest studies appeared in the 1980s and many additional reports have

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confirmed and extended this general finding during the past 25 years [2–5]. Initially, research was directed towards tumour cell lines such as breast and colon cancer for which epidemiological data suggested a link between dietary fat intake and cancer incidence. Over time, many other tumour types were found to be sensitive to the inhibitory effects of fatty acids, and relationships were identified among the number of carbon atoms and the number of double bonds present in the fatty acids and their respective tumour inhibitory properties. The position of the double bonds within the fatty acid carbon chain was also found to cause differences in the effects on tumour cell proliferation [5–8]. While the n-3 PUFAs had strong inhibitory effects on tumour cell proliferation, the n-6 PUFA gamma-linolenic acid (GLA) was often found to be one of the most inhibitory and most selective fatty acids in studies comparing tumour cells with normal cells *in vitro* [5, 9]. The susceptibility of individual fatty acids to peroxidation was also identified as an important factor in their ability to inhibit tumour cell proliferation [4–6, 9–13].

The epidemiological data have made a strong case for an increased risk of cancer development in populations consuming occidental diets rich in saturated fatty acids. In addition to the saturated nature of the majority of the fatty acids, the small quantity of PUFAs present in the diet is typically of the n-6 family. Thus, the relative paucity of dietary n-3 PUFAs versus n-6 PUFAs has been advocated as one of the risk factors for cancer development. Further research has pointed to changes in breast cancer incidence in populations with traditionally high dietary ingestion of n-3 PUFAs when members of the population have moved to other regions of the world and adopted local dietary habits [14–17].

In vivo studies have also shown significant effects of PUFAs upon tumour development and progression in animal models. Many of the animal models have followed (1) tumour development after exposure to carcinogenic stimuli, (2) progression of established tumours or (3) development, progression and eventual metastasis of tumours, each in the presence of different types of PUFA. Early studies in mouse models of tumour growth found that supplementation of an essential fatty acid-deficient diet with the n-6 PUFA linoleic acid (LA) stimulated tumour growth. In contrast, the n-6 PUFA arachidonic acid (AA) had no significant effect on tumour growth rates. Most of the animal studies have classified the n-6 PUFA LA as a stimulator of tumour growth in *in vivo* models [18–21].

The picture is quite different when animal tumours are exposed to n-3 PUFAs, with most studies finding marked inhibition of tumour growth in the presence of these PUFAs. Initial work in this field tested the effects of n-3 PUFAs in the complex mixtures of PUFAs found in fish oils. Hepatoma growth was significantly reduced when

animals were fed diets enriched with fish oil. Similar findings were reported for breast and prostate tumour growth rates [22–28]. In a mammary tumour model in mice dietary supplementation with a fish oil, supplement containing 12% docosahexaenoic acid (DHA) and 18% eicosapentaenoic acid (EPA) also reduced tumour growth. More recent protocols have used highly purified EPA or DHA in models of colon, breast and prostate tumour development after exposure to carcinogenic compounds [14, 25]. In both cases, the n-3 PUFAs significantly reduced the incidence and development of tumours in these models. The importance of fatty acids during metastatic spread has also been highlighted in studies where n-3 PUFAs have been found to decrease the number of metastatic foci in a breast cancer model [26]. In brain tumours, studies comparing dietary fat intake and brain tumour incidence are scarce and have not found convincing epidemiological evidence of a significant association between, for example, gliomas and dietary PUFAs [29].

Effects of Fatty Acids on Mitochondrial Metabolism

The utilisation of fatty acids for energy production via the mitochondrial beta-oxidation pathway depends upon the availability of fatty acyl CoAs within the mitochondrial matrix. The provision of fatty acyl CoAs for beta-oxidation relies on the concerted activity of carnitine palmitoyltransferase I (CPT I), the carnitine-acylcarnitine translocase (CACT) and carnitine palmitoyltransferase II (CPT II) in order to pass from the cytoplasm to the mitochondrial matrix. In tumour cells, fatty acid oxidation rates are often reduced in comparison with the normal tissue of origin, and it is, therefore, often stated in the literature that tumour cells do not oxidise fatty acids. This idea is propagated by the lower quantity of mitochondria often found in tumour versus normal tissue and by the common tendency of tumour cells to utilise glucose as their principal energy source via the glycolytic pathway [30, 31]. However, the evidence for a lack of fatty acid oxidation by tumour cells was scarce, and when CPT activities and fatty acid oxidation rates of various tumour cells were analysed, it became apparent that many different tumour types do in fact retain fatty acid oxidative capacity, which may be utilised during specific periods of tumour growth [32–35]. The tumour cell's ability to oxidise fatty acids was found to vary considerably depending on the fatty acid, which is related to the specificity of CPT I for substrate [35, 36].

With this in mind, studies in the author's laboratory tested the effects of PUFAs on the capacity of tumour cells to oxidise fatty acids and found significant effects of several fatty acids. Both GLA and AA caused approximately 60% inhibition of tumour cell CPT I activity, while

oleic and LA had no effect on CPT I activity. Both GLA and AA also caused approximately 50% inhibition of [$1\text{-}^{14}\text{C}$]-palmitate oxidation to $^{14}\text{CO}_2$ [37]. These effects were blocked by the inhibition of cyclooxygenase with indomethacin suggesting a role for prostaglandins (PGs) in this effect. Indeed, addition of PGE_1 and PGE_2 to the tumour cells also caused a significant inhibition of both CPT I activity and [$1\text{-}^{14}\text{C}$]-palmitate oxidation to $^{14}\text{CO}_2$. The alterations in fatty acid oxidation within the tumour cells could potentially cause disturbances in other pathways of fatty acid metabolism including triacylglycerol and cholesterol synthesis, membrane biosynthesis and eicosanoid synthesis amongst others. These changes in fatty acid metabolism could lead to selective inhibition of tumour cell proliferation, and in the cell lines tested, this was found to be the case. In the same study, a potential positive control was used in the form of the colon cancer cell-line HT29 whose proliferation was known to be unaltered by PGE_2 [38]. Supporting the hypothesis that altered CPT I could result in altered proliferation while lack of inhibition would result in unchanged proliferation, the HT29 cells exposed to GLA, AA, PGE_1 or PGE_2 showed no alterations in CPT I activity and no alterations in proliferation [37]. Later reports found that HT29 DNA fragmentation, indicating apoptosis, was unaffected by various PUFAs with the exception of DHA, which was a strong inhibitor of cell proliferation in this cell line [39].

In vivo studies using the Walker 256 carcinosarcoma model in the rat reported that modification of the fatty composition of the diet with n-3 fatty acids caused significant changes in mitochondrial fatty acid composition. The activity of the mitochondrial acyl CoA synthase was also significantly increased in the tumours from the PUFA-rich diet [36, 40]. These changes in mitochondrial metabolism were accompanied by a significant decrease in mitochondrial membrane potential (MMP). Further studies using the same animal model found that GLA also caused a marked increase in mitochondrial acyl CoA synthase activity, and again, MMP was decreased by the PUFA diet [40].

A similar 5.5% GLA diet was subsequently found to cause a 61% inhibition of CPT I activity in the Walker 256 carcinosarcoma, which was accompanied by both a decrease in the malonyl CoA sensitivity of the enzyme and a decrease in its affinity for palmitoyl CoA substrate [36]. The separation of the mitochondria into outer membrane, contact sites and inner membrane fractions provided detailed information on the variations in susceptibility of individual biological membranes to dietary alterations in fatty acid composition. The outer membrane was highly susceptible to dietary influence on fatty acid composition, with the contact sites less so and the inner membrane fraction more refractory to changes enforced by the diet.

Another important change that was found in these treated tumours was the large decrease in mitochondrially bound hexokinase (HK) activity present in the tumours from 67% (control tumour) to 39% (GLA tumour) of the total detectable HK activity [36]. Despite this large change in localisation, the total HK activity was itself unchanged. Tumour cells very often have a large part of their HK I and/or II enzyme bound to the mitochondria, which is believed to improve energy production by placing the first step of glycolysis very close to the mitochondrial source of ATP [41–43]. Such a change in the intracellular localisation of this important enzyme could lead to a situation where intramitochondrial ATP synthesis becomes less efficiently coupled to cytoplasmic glucose-6-phosphate generation. In turn, this reduced efficiency could alter the rate of glycolysis and alter the energy balance within the rapidly proliferating tumour cells. This study provided evidence that fatty acids could influence the capacity of HK to bind to the mitochondrial outer membrane protein voltage dependent-anion channel (VDAC). Long-chain fatty acids have been reported to inhibit heart HK (type I HK), which is also known to bind to mitochondria [44]. It was proposed that high intracellular concentrations of fatty acids may serve as an additional control over HK binding and function in the heart, and this is a possibility that may be extrapolated to tumour HK in some cases where high-intracellular fatty acid concentrations may be achieved. The loss of bound HK from the VDAC has recently been directly related to loss of protection from apoptosis provided by this HK–VDAC interaction [45]. The capacity of HK I and HK II to interact with VDAC points to the role of these proteins in promoting tumour cell survival through binding to VDAC, which in turn inhibits the release of cytochrome *c* from the mitochondrial intermembrane space [42, 43, 46, 47]. These recent reports on HK–VDAC interaction and apoptosis further support the previous data from our laboratory linking loss of HK from mitochondria as having an important role in the subsequent loss of MMP and apoptosis described later in this review [48–52].

The susceptibility of individual fatty acids to peroxidation was previously identified as an important factor in their ability to inhibit tumour cell proliferation [5, 6, 9–12]. Studies in the Walker 256 rat carcinosarcoma cell line and in bladder tumour cells found that GLA or EPA caused significantly increased thiobarbituric acid reactive substances (TBARS) formation, indicative of lipid peroxidation, and also increased oxidation of dihydroethidium, indicative of ROS generation such as superoxide [40, 48]. The superoxide and lipid peroxide production pointed to altered mitochondrial function [53], and when mitochondrial enzyme activities were determined, both NADH cytochrome *c* oxidoreductase (complex I + III) and cytochrome *c* oxidase (complex IV) activities were greatly reduced by the presence

of PUFAs [49]. Together with reduced electron transport capacity, the mitochondrial fraction was seen to lose its cytochrome *c* content in a controlled manner, which did not involve generalised mitochondrial rupture [49]. The loss of cytochrome *c* would directly alter the mitochondrial ability to generate ATP and in addition serves as a stimulus for cell death. The n-6 PUFA AA has been reported to uncouple mitochondrial respiration by the selective inhibition of complex I and complex III and to cause an increase in ROS production by the mitochondria [54].

The reduced mitochondrial function was accompanied by an increase in glucose utilisation and lactate production by the tumour cells [49, 55]. This change in metabolic activity would compensate to some extent the oxidative deficit produced by the presence of PUFAs and was reflected in the reduction of intracellular ATP by between 205 and 27% depending on the PUFA used. Despite metabolic adaptations made by the tumour cells, PUFAs led to reduced MMP suggestive of the opening of the mitochondrial permeability transition (MPT) pore [47, 52, 53]. Oxidative stress was reported to directly influence the occurrence of MPT and cytochrome *c* release [56–58]. It has been proposed that when mitochondrial complex I or II activity is inhibited, cancer cells may compensate with an increased flux through glycolysis to generate sufficient ATP to sustain the apoptotic pathway and cause cell death [59].

The increased acyl CoA found in the PUFA exposed cells may also be relevant to the inhibition of mitochondrial metabolism and the induction of apoptosis, as acyl CoAs are known to inhibit the mitochondrial adenine nucleotide transporter (ANT). This protein is present at contact sites and is one of the components of the MPT pore [47, 52, 53]. It is possible that increased intracellular acyl CoA concentrations in the microenvironment of the mitochondrial membrane, due to increased substrate availability and increased enzyme activity [40, 48], could inhibit the ANT and compromise energy metabolism. While this has not been directly proven, it is an important factor that could drastically alter mitochondrial energy metabolism and organelle integrity [47]. A significant reduction in mitochondrial oxidation would also lead to a reduction in citrate production and efflux from the mitochondria, which would interfere directly with the activity of ATP-citrate lyase and fatty acid synthesis in the rapidly proliferating tumour cells [46, 60].

Interestingly, in HT29 colon, tumour cells LA and AA caused modifications in total phospholipid and mitochondrial cardiolipin fractions, which were accompanied by decreases in MMP [61]. In Morris hepatoma cells, cardiolipin composition was significantly altered by LA, while in Walker 256 cells, GLA and EPA altered cardiolipin composition [49]. EPA also significantly altered the cardiolipin fatty acyl chain composition in bladder tumour

cells [55, 62], and in both the Walker 256 cells and bladder tumour cells, MMP was significantly reduced. The peroxidation of cardiolipin has been proposed to be an initial event in the release of cytochrome *c* from the mitochondria, and enriching cardiolipin with PUFA acyl chains may lead to increased peroxidation and will be addressed later in this review [55, 63].

Fatty Acids and Mitochondrial Ultrastructure

Alterations in the fatty acid composition of the diet are known to influence the mitochondrial composition and their metabolism as described in the previous section. These findings raised the question of whether changes in mitochondrial metabolism may be related to actual ultrastructural changes induced by PUFAs. Initial studies found that in the Walker 256 carcinosarcoma model, a high-GLA-containing diet caused an increase in mitochondrial area and volume, while a low-GLA-containing diet caused no difference from a control diet [48]. These studies were extended in order to investigate the relationship between the alterations in mitochondrial metabolism and the organelle ultrastructure in vivo. Particular attention was paid to alterations in ultrastructure previously considered to be related to alterations in intracellular metabolic state, including the quantity of cytoplasmic lipid droplets, the number of mitochondrial matrix granules and the number of mitochondrial membrane contact sites [36]. A strong relationship has been proposed among the number of contact sites and matrix granules and the metabolic activity of the mitochondria [64, 65].

Quantitative biochemical analysis of the incorporation of GLA into various lipid fractions found that while neither phospholipid nor cholesterol fractions were altered in quantity, the triacylglycerol fraction almost doubled in size in the GLA dietary group [36]. Ultrastructural analysis of the lipid droplet fraction confirmed the biochemical data showing a doubling of the number of lipid droplets per total cell area and a doubling of the lipid droplet area per unit cell area. In this study, the mitochondrial area and volume were also increased by GLA as previously found in [48]. However, the number of mitochondria per unit cell area remained unchanged indicating that GLA did not cause mitochondrial proliferation. GLA exposure caused a significant reduction in the number of matrix granules to approximately half of those present in the control tumour. The surface density of mitochondrial cristae was reduced to almost half that of the control tumour, which would in itself reduce the physical space available for membrane insertion of the enzymes involved in the respiratory chain and may also in part explain the reduction in complex I, III and IV activity in PUFA-treated tumour cells [36, 49]. Changes in

cristae structure have been linked to changes in cytochrome *c* localisation within the mitochondria from the cristae to the intermembrane space, which may facilitate cytochrome *c* release during the induction of cell death [47].

This finding was accompanied by a 66% decrease in mitochondrial membrane contact sites in the GLA-exposed tumours [36]. Such a large difference in contact site number may have a dramatic effect on mitochondrial function as many tumours are already depleted of contact sites when compared with their normal tissues of origin [66]. Thus, a further reduction caused by GLA exposure may push the mitochondria to the limit of their capacity to function with insufficient contact sites available for essential metabolic activities. In earlier studies, fatty acids were also linked to a decrease in the formation of contact sites, while essential fatty acid deficiency can produce megamitochondria, proving that fatty acids can significantly alter mitochondrial ultrastructure [64, 67].

The large change in contact site number may be directly related to the significant decrease in CPT I activity previously mentioned, as CPT I activity, at least in rat liver, has been found to have differing kinetic properties when located at the outer mitochondrial membrane or at contact sites [68–70]. Taking into consideration the large decrease in contact sites in the GLA-exposed tumours, most of the CPT I enzyme was relocated to the outer membrane fraction rather than the contact site fraction. This change in localisation would mean that CPT I would (1) be exposed to a different fatty acyl chain composition in the outer membrane in comparison with the composition of the contact sites and (2) be exposed to an altered outer membrane composition in the GLA diet in comparison with the control diet outer membrane. Thus, the loss of contact sites causes a direct effect on the enzyme by altering its membrane environment through both altered location and altered compositions.

Fatty Acid Induction of Cell Death

In vivo studies in the Walker 256 carcinosarcoma model found that cell cycle changes occurred in the presence of n-3 PUFA-rich diets leading to a decrease in cells in the G1 phase and an increase in cells in the G2-M phase [40]. As previously mentioned, these changes were associated with an increase in TBARS content in the tumour cells, and a large decrease in cell proliferation and an increase in the apoptotic index were also found. Interestingly, in this model, EPA caused a higher rate of TBARS production along with a greater inhibition of tumour growth in vivo than the more unsaturated DHA. Both GLA and AA inhibited the cell cycle in various other tumour cells causing reductions in S-phase and induction of apoptosis

[71–73]. DHA has also been reported to exert similar effects in leukaemia cells with S-phase cycle arrest, increased p21 expression and increased pro-caspase-3 cleavage leading to increased apoptotic rate [74, 75].

Further studies using the same Walker 256 tumour model found that GLA produced a dose-dependent inhibition of tumour growth with an almost 50% inhibition in the high-GLA group, which consumed a diet containing 5.5% GLA. Once again, the apoptotic index within the tumour tissue was significantly increased by the PUFA-rich diet [48]. Corroborating the fact that more cells were apoptotic, the nuclear size and shape factor of GLA exposed tumours was significantly reduced as would be expected in cells with pyknotic and apoptotic nuclei present [36].

As previously mentioned, a significant change in the intracellular localisation of mitochondrially bound HK could lead to a situation where intramitochondrial ATP synthesis becomes less efficiently coupled to cytoplasmic glucose-6-phosphate generation. In turn, this reduced efficiency could alter the rate of glycolysis and alter the energy balance within the rapidly proliferating tumour cells, thereby producing a signal for the induction of apoptosis due to reduced ATP production. The decreased MMP found in GLA-exposed tumours may be related to changes in the MPT pore induced by the alterations in mitochondrial HK binding [76–79]. In a recent study, mitochondrial pore-targeted drugs caused MPT and subsequently apoptosis in temozolamide-resistant glioma cells [79].

An article published around the time of Colquhoun [36] reported that mitochondrially bound HK was a modulator of early apoptotic events through the serine–threonine protein kinase, PKB/Akt [76]. Akt stimulates the binding of HK to the mitochondria, thereby increasing the capacity for coupling mitochondrial ATP synthesis to the first step of the glycolytic pathway catalysed by HK [46]. Recent work on Akt has shown that the Akt protein has an Achilles' heel. Normally, Akt is able to inhibit apoptosis induced by many different apoptotic stimuli. However, Akt is unable to protect against ROS-mediated apoptosis and actually sensitises cells to this form of cell death [80]. Tumour cells overexpressing Akt activity, often due to PTEN mutation, may therefore be good targets for chemotherapeutic agents that produce ROS, which would permit the exploitation of this aspect of Akt to increase the tumour cell kill rate. The increase in ROS typically present in PUFA-exposed tumour cells could be particularly effective in tumours overexpressing Akt.

Another finding that suggests a link between metabolic changes in fatty acid-treated tumour cells and the induction of apoptosis comes from the previously mentioned studies involving CPT I. The suggestion that CPT I could be involved in apoptosis was originally proposed by Paumen et al. [81, 82]. These data showed the capacity of the anti-

apoptotic protein Bcl2 to bind to CPT I in a yeast two-hybrid system. In interleukin-3-dependent cells undergoing apoptosis after interleukin-3 deprivation, CPT I expression was increased. This article raised the possibility that an increase in CPT I offered protection from palmitate-induced apoptosis by improving the clearance of long-chain acyl CoAs from the cytoplasm [81, 82]. In cardiomyocytes, a reduction in CPT I in the presence of high concentrations of palmitate was associated with a decrease in MMP and an increase in apoptosis [83]. Studies in the author's laboratory found that PUFAs caused inhibition of CPT I, inhibition of fatty acid oxidation and increased cytoplasmic acyl CoA and triacylglycerol content in both in vitro and in vivo tumour models and identified a strong correlation among the inhibition of CPT I, the inhibition of cell proliferation and the induction of apoptosis in a human larynx tumour cell line [84]. In this cell line, the PUFA effects on CPT I activity found at 6 hours after exposure were correlated with the apoptotic events seen at 12 and 24 hours after exposure, suggesting that the inhibition of mitochondrial CPT I and fatty acid oxidation may lead to the induction of apoptosis in certain tumour cells.

The altered composition of cardiolipin found in several tumour cells exposed to PUFAs has been proposed to increase its likelihood of suffering peroxidation [49, 55]. Recent studies have shown that cardiolipin peroxidation is an important initiating factor in the induction of some forms of cell death [59, 85]. The release of cytochrome *c* from the mitochondria in PUFA-treated cells involves the peroxidation of cardiolipin in order to release cytochrome *c* after which apoptosis can be induced via the remainder of the intrinsic mitochondrial pathway. The increased ROS typically found in PUFA-treated cells would tend to increase the probability of oxidising cardiolipin. In PUFA-treated Walker 256 cells, bladder tumour cells and larynx tumour cells caspase-3 were activated after cytochrome *c* release from the mitochondria, and this was followed by DNA fragmentation readily detected by in situ TUNEL - terminal deoxynucleotidyl transferase mediated dUTP nick-end labelling assays and FACS - fluorescence-activated cell sorting analysis. These changes were accompanied by significant loss of MMP of up to 50% of control cell values [49, 55]. Interestingly, complexes I, III, IV and V of the mitochondrial respiratory chain require cardiolipin for optimal function; complexes III, IV and V contain cardiolipin in their quaternary structure and cardiolipin is also required by several mitochondrial substrate carriers including the phosphate carrier, the ANT and the CACT [85]. The changes found in cardiolipin fatty acyl composition, which increase its tendency for peroxidation, could lead to deficiencies in functional cardiolipin within the tumour cell and may in part explain the reductions in mitochondrial respiratory capacity. The involvement of cardiolipin with the CACT is intriguing as cardiolipin is also associated with changes in CPT I activity [86]. It is possible that increased

peroxidation of cardiolipin in the PUFA-treated tumour cells is partially responsible for the reduction in CPT I activity and fatty acid oxidation previously mentioned [36, 37]. Recent studies have also raised the question of whether interactions between the pro-apoptotic Bcl2 family members and peroxidised cardiolipin may regulate and/or stimulate cytochrome *c* release and suggest that the manipulation of cardiolipin peroxidation may be a target for the sensitisation of tumour cells to apoptosis [85]. For further reading on PUFAs and their ability to induce apoptosis, an excellent recent review is recommended [87].

Implications for Neuro-oncology

Two of the earliest studies linking fatty acids to inhibition of tumour cell growth involved neuroblastoma cells and glioma cells [4, 88]. Since then, the n-6 PUFAs GLA and AA have been shown to inhibit glioma cell proliferation, inducing ROS production and leading to cell death [89]. Brain tumours are often deficient in essential fatty acids containing far more saturated than unsaturated fatty acids in their cell membranes, which may reduce their intracellular production of ROS [13, 90, 91]. In a group of human gliomas including oligoastrocytomas, anaplastic astrocytomas and glioblastoma multiforme, AA and GLA were found to stimulate tumour cell peroxidation and apoptosis [89, 92, 93]. Normal brain cells were less susceptible to ROS production and apoptosis in the presence of PUFAs, and necrotic subpopulations of tumour cells also produced less ROS than other regions of the tumour tissue.

Several human studies have tested the effects of GLA on glioma progression, and although the design of these studies was less than ideal, the data suggest that GLA has inhibitory effects on glioma progression in vivo. In these studies, doses of up to 1 mg GLA/day were used for up to 20 days, and the dosage was well tolerated in the majority of patients, some of which showed signs of disease stabilisation during the studies [94–98].

Further in vitro studies have shown that GLA significantly inhibited glioma cell proliferation in three-dimensional spheroids grown in collagen gels and stimulated a dose-dependent increase in apoptosis in the same model [99]. In the C6 cell line, both GLA and EPA were reported to inhibit cell migration on a plastic substrate, while DHA was found to have no effect at similar concentrations [100] (unpublished data). Although migratory tumour cells are considered to be non-proliferative, glioma cells can rapidly stop migration, proceed through cell division and enter into a migratory phase again [101]. With this in mind, mitosis was evaluated in the migratory field and found to be significantly reduced in the presence of either GLA or EPA [100].

More recent studies in the author's laboratory have attempted to identify some of the molecular mechanisms involved in the effects of GLA upon glioma cells both in vitro and in vivo. The in vitro studies found that, as expected, GLA caused a significant decrease in cell proliferation. From the experimental findings, it was apparent that the reduced proliferative rate was accompanied by a reduction in the expression of the protein E2F1, which is required for the entry of cells into the S-phase of the cell cycle and by a reduction in the number of cells in the S-phase when analysed by FACS [102]. In HT29 colon tumour cells, DHA reduced the ability of E2F1 to bind to DNA suggesting reduced activation of E2F1 by DHA and resulted in reduced S-phase [103]. The effects of GLA could only be partially reverted by the natural antioxidant alpha-tocopherol suggesting that ROS production is only partially responsible for GLA-induced changes in cell proliferation in the C6 glioma cell in vitro [102].

GLA caused significant alterations in the expression of the pro-apoptotic protein Bax and the anti-apoptotic protein Bcl2, which are both involved in the control of the intrinsic mitochondrial apoptosis pathway [51, 104, 105]. However, at the concentration of GLA used in this study, C6 cells did not enter into apoptosis even though MMP is reduced [100, 102]. However, at higher concentrations, GLA does induce cell death in these cells (unpublished data). Although the increase in Bax expression would suggest a possible increase in apoptosis [53], in neuroblastoma cells Bax can associate with the DNA repair protein Ku70, thereby blocking Bax's pro-apoptotic function [106, 107]. Ku70 can also function as a physiological inhibitor of Bax-induced apoptosis. Ku70/Ku80 heterodimers represent the regulatory subunit of the DNA-dependent protein kinase (DNA-PKc) which plays an important role in DNA damage repair. In GLA-treated C6 cells, Ku80 expression is significantly reduced which may leave more Ku70 protein free in the cytoplasm where it could bind to Bax and annul its potentially pro-apoptotic effects. Interestingly, Ku proteins are related to hyperproliferation in tumour cells, and inhibition of their expression may lead to decreased cell proliferation as they are implicated in the maintenance of the proliferating cell nuclear antigen (PCNA) on chromatin and can associate with E2F1 [107–109]. The oxidative stress produced by GLA may also be responsible for the reduced Ku80 levels, as oxidative stress in other studies was reported to cause nuclear loss of Ku proteins [110]. The large reduction in Ku80 protein expression could be directly related to impaired DNA replication resulting in cell cycle changes and the micronucleus formation seen in these GLA-exposed cells. Indeed RNAi studies blocking production of Ku80 were reported to cause cell cycle block in G1, reduced S-phase and decreased PCNA expression [111]. GLA was shown to reduce PCNA expression in

HeLa cells, and this was associated with possible loss of DNA repair capacity [73]. Recent studies have shown that blocking or reduction of Ku protein function can increase tumour cell sensitivity to both radiotherapy and chemotherapy [112–114].

GLA and other PUFAs are known to increase the sensitivity of tumour cells to both radiotherapy and chemotherapy, although the mechanisms behind these effects are poorly understood [9, 115–119]. Thus, the fact that GLA alters DNA repair protein Ku80 expression in glioma cells may be involved in its capacity to improve tumour cell responses to chemotherapy and radiotherapy and even improve the multi-drug resistance of tumour cells [120, 121]. Of relevance to this is the finding that in C6 glioma cells, the improved sensitivity to radiotherapy did not appear to be directly related to increased ROS generation [121]. It is also possible that changes in energy metabolism and ATP synthesis may impede the activity of ATP-dependent multi-drug resistant proteins so often active in tumour cells [122].

The in vivo studies from Leaver et al. [121, 123] have tested the efficacy of varying doses of GLA applied directly into the tumour bed via osmotic pumps using the orthotopic C6 glioma model. Normal brain morphology was well preserved with no difference in reactive changes at the infusion site between the control and GLA-treated tumours. In tumours exposed to 2mM GLA for 7 days of continuous infusion at 1 μ l/h, the tumour size was less than 50% of the phosphate-buffered saline control-treated tumours and the Ki67 labelling index in the GLA-treated tumours was significantly decreased [123].

In a recently published study from the author's laboratory, GLA was applied directly into the tumour bed via osmotic pumps using the same orthotopic C6 glioma model used by Leaver et al. [121, 123, 124]. The infusion rate was reduced to 0.5 μ l/h, and the treatment consisted of 5 mM GLA infusion for a total of 14 days with the expectation that a higher dose for an extended period of time would improve tumour response to GLA [124]. The average tumour area of the GLA treated tumours was reduced by 75% in comparison with the control cerebrospinal fluid-treated tumours, and this was accompanied by a change in tumour fatty acid composition. The expression of proteins involved in cell cycle control and apoptosis were significantly changed by GLA treatment including cyclin D1, p53 and p27 expression, and these changes were concomitant with decreased cell proliferation, decreased bromodeoxyuridine incorporation in the S-phase of the cell cycle and increased apoptosis. The possible role of increased p53 expression in altered mitochondrial energy metabolism should also be taken into consideration in this in vivo model as another potential mechanism for reducing cell proliferation [60, 125–127].

Together with the changes in cell proliferation and apoptosis, GLA was found to decrease the expression of proteins involved in angiogenesis including vascular endothelial growth factor (VEGF), ERK1/2, VEGF receptor Flt1 and the matrix metalloproteinase MMP2 and to decrease the microvessel density of the tumour. The findings from this particular study point to several areas in which GLA exerts either direct or indirect effects on tumour growth [124]. Pancreatic tumour cells treated with DHA were also found to have reduced ERK1/2 expression along with increased expression of p27, and in mammary tumours, DHA caused a decrease in proteosomal degradation of p27 [128, 129]. Exposure of HT29 colon cancer tumours to n-3 PUFAs in vivo also caused decreased expression of VEGF, decreased ERK1/2 phosphorylation and decreased microvessel density, leading to decreased tumour growth [130].

Particularly important in these glioma studies is the identification of a dual action of GLA both inhibiting tumour cell proliferation and angiogenesis in vivo [123, 124]. Previous studies reported that GLA altered the expression of proteins involved in endothelial junction formation such as VE-cadherin and occludin and interfered with endothelial tube formation [131–133].

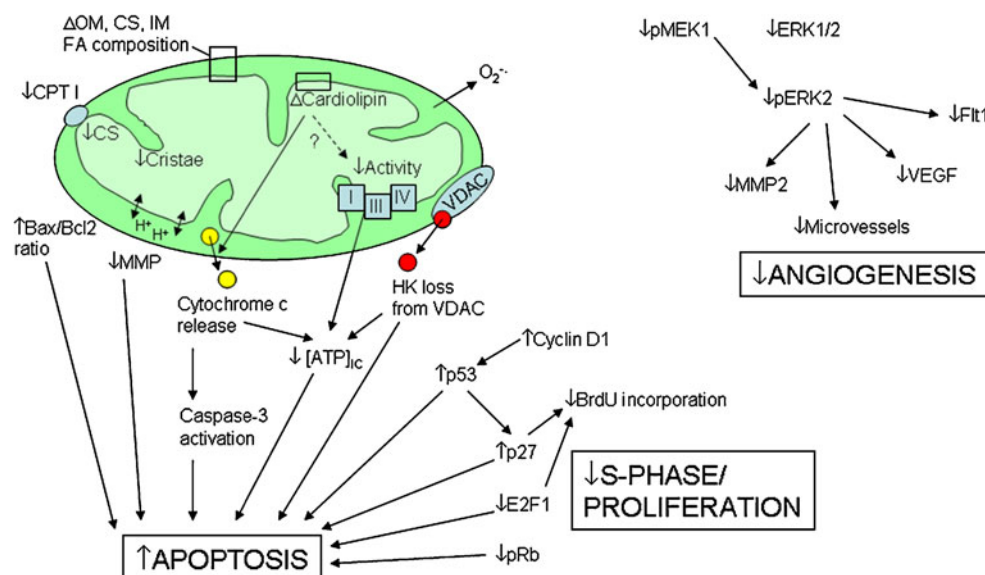
A better understanding of the metabolism of fatty acids and eicosanoids in brain tumours and their influence on energy balance will be fundamental to the possible targeting of mitochondria in tumour treatment [134–136]. A recent review article stated that “the mitocan class of drugs includes compounds that affects such mitochondria-associated activities as hexokinase inhibition, activation of the mitochondrial permeability transition pore, inhibition of bcl2 anti-apoptotic proteins and blocking of the electron transport/ respiratory chain” [59]. Through the author’s own and many other studies, GLA can readily be placed in this

category, and it is reasonable to suggest that the use of combination therapies with complimentary targets and modes of action together with PUFAs such as GLA may lead to gains in treatment efficacy in gliomas. Fig. 1 contains a summary of the ideas and data presented in this review.

Conclusions and Future Directions

It is clear from the data reviewed in the preceding sections that PUFAs exert significant inhibitory effects on tumour cell proliferation. This is in part related to the increased production of ROS, which occurs when the tumour cells are exposed to PUFAs. The role of CPT I in the function of normal astrocytes is poorly understood and even less is known of its importance in gliomas, leaving many questions unanswered regarding PUFA effects on fatty acid oxidation and eicosanoid metabolism in gliomas and is an area that deserves investigation [136–140]. HK binding to mitochondria and its involvement in the inhibition of apoptosis through interactions with VDAC, as well as the possibility of inhibiting ANT through increased acyl CoA concentrations, are also areas of interest for future glioma research [141–143]. The question remains whether PUFAs are able to alter glioma HK binding to mitochondria and whether this may be an important target for exploitation when taking into account the additional involvement of HK in the activity of Akt, a protein often activated in gliomas due to mutation of PTEN. The ability of PUFAs to increase tumour sensitivity to both radiotherapy and chemotherapy also merits further attention in gliomas, as it may provide a means to improve tumour response to conventional treatment options. Research into the targeting of mitochondrial

Fig. 1 Effects of PUFAs on tumour metabolism. OM outer mitochondrial membrane; CS mitochondrial membrane contact site; IM mitochondrial inner membrane; FA fatty acid; CPT I carnitine palmitoyltransferase I; MMP mitochondrial membrane potential; I, III, IV mitochondrial respiratory complexes I, III and IV; VDAC voltage-dependent anion channel; HK; BrdU bromodeoxyuridine; MMP2 matrix metalloprotease 2; VEGF vascular endothelial growth factor; Flt1 VEGF receptor Flt1; $O_2^{\cdot-}$ superoxide



metabolism in an attempt to direct tumour cells towards apoptotic and, more recently, autophagic cell death is gaining momentum in the literature and may provide novel strategies for the treatment of primary brain tumours such as gliomas [51, 52, 59, 79, 135, 144, 145].

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