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PPARδ prevents radiation-induced proinflammatory responses in microglia via transrepression of NF-κB and inhibition of the PKCα/MEK1/2/ERK1/2/AP-1 pathway

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Abstract

Partial or whole-brain irradiation is often required to treat both primary and metastatic brain cancer. Radiation-induced normal tissue injury, including progressive cognitive impairment, however, can significantly affect the well-being of the approximately 200,000 patients who receive these treatments each year in the US. Although the exact mechanisms underlying radiation-induced late effects remain unclear, oxidative stress and inflammation are thought to play a critical role. Microglia are key mediators of neuroinflammation. Peroxisomal proliferatoractivated receptor (PPAR)δ has been shown to be a potent regulator of anti-inflammatory responses. Thus, we hypothesized that PPARδ activation would modulate the radiation-induced inflammatory response in microglia. Incubating BV-2 murine microglial cells with the PPARδ agonist, L-165041, prevented the radiation-induced increase in: i) intracellular reactive oxygen species generation, ii) Cox-2 and MCP-1 expression, and iii) $IL-1\beta$ and TNF- α message levels. This occured, in part, through PPARδ-mediated modulation of stress activated kinases and proinflammatory transcription factors. PPARδ inhibited NF-κB via transrepression by physically interacting with the p65 subunit, and prevented activation of the PKCα/MEK1/2/ERK1/2/AP-1 pathway by inhibiting the radiation-induced increase in intracellular reactive oxygen species generation. These data support the hypothesis that PPARδ activation can modulate radiationinduced oxidative stress and inflammatory responses in microglia.

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Keywords

Ionizing radiation; PPARδ; radiation-induced brain injury; microglia; inflammation; NF-κB; PKCα/MEK1/2/ERK1/2/AP-1 pathway

> Each year, approximately 200,000 patients will receive fractionated partial or whole-brain irradiation (fWBI) as treatment for primary or metastatic brain cancer [1]. Unfortunately, the radiation dose that can be delivered to the tumor is limited by the risk of toxicity to the surrounding normal brain tissue. Radiation-induced brain injury can lead to cognitive deficits several months to years after irradiation that affect patients' quality of life (QOL) [2– 4]. This diminished QOL has become an important concern for these long-term survivors of brain irradiation, and is recognized as one of the most important measurements of brain tumor therapy outcomes in clinical trials, second only to survival [5, 6]. Although clinical trials have demonstrated that short-term interventions can modulate cognitive impairment [7], there are no proven long-term treatments for radiation-induced cognitive deficits; therefore, it is important to investigate new therapeutic approaches [8].

> Although it was once believed that the brain was not susceptible to inflammation, studies have now demonstrated that inflammatory responses do occur and may contribute to radiation-induced brain injury. In vivo studies indicate that there is an increase in proinflammatory mediators within hours of irradiating the rodent brain [9–11]. Microglia are considered to be one of the key mediators of neuroinflammation [12–15]. In an uninjured brain, ramified microglia actively monitor the microenvironment to ensure that the brain is maintaining homeostasis [12]. Following injury, microglia become activated, a process characterized by rounding of the cell body, retraction of cell processes, proliferation, and an increased production of cytokines, chemokines, and reactive oxygen species (ROS) [12–15]. Although microglial activation plays an important role in phagocytosis of dead cells, sustained activation is thought to contribute to a chronic proinflammatory state in the brain [13, 15]. In vivo studies in rodents indicate that radiation leads to an increase in microglial activation [16, 17]. Irradiating microglia cells in vitro leads to an increase in proinflammatory mediators, such as the cytokines TNF- α and IL-1 β , and the chemokines MCP-1 and ICAM-1 [18–20].

> Radiation-induced chronic oxidative stress and inflammatory responses produced by microglia may: i) lead to a decrease in neurogenesis in the hippocampus, a critical region for learning and memory; and/or ii) alter the environment of the neurogenic niche and, in turn, the functions of the pre-existing neurons [15, 16, 21]. Studies in rodents demonstrate that the administration of anti-inflammatory drugs can decrease radiation-induced microglial activation. This decrease has been associated with an improvement in hippocampal neurogenesis [16, 17]. Moreover, administration of eicosapentaenoic acid, a polyunsaturated fatty acid with anti-inflammatory properties, restored the altered long-term potentiation (LTP) of hippocampal slices following irradiation of the rat brain [21]. LTP is thought to underlie memory and cognitive function; it is a measure of signal transmission between two neurons [22]. These findings provide a strong rationale for investigating anti-inflammatory therapies to mitigate radiation-induced brain injury.

PPARδ is a member of the nuclear hormone receptor superfamily of ligand-activated transcription factors and one of three subtypes $(α, δ, and γ)$ that make up the PPAR family [23, 24]. PPARs regulate transcription by heterodimerizing with the retinoid \times receptor (RXR) and binding to PPAR response elements (PPREs). PPREs, located in the enhancer regions of genes, consist of an AGGTCA hexameric direct repeat separated by one or two nucleotides [25]. PPAR subtypes are encoded by different genes, have unique tissue

distributions, and exhibit overlapping and differential functions. PPARδ is thought to be expressed ubiquitously; it is the predominant PPAR subtype in the CNS [26]. Several studies have demonstrated that PPARδ activation can modulate oxidative stress and inflammatory processes [27–29]. PPARδ mediates many of its anti-inflammatory effects by preventing activation of stress-activated kinases and proinflammatory transcription factors [30, 31]. In addition, PPARδ has been shown to regulate oxidative stress by activating transcription of antioxidant genes, such as catalase and superoxide dismutase [32, 33].

Recent studies suggest that PPARδ agonists may ameliorate the severity of various acute and chronic CNS pathologies, including stroke, multiple sclerosis, and Alzheimer's disease, in large part, by modulating the oxidative stress and proinflammatory responses associated with these diseases [34–36]. The role of PPARδ in the modulation of radiation-induced brain injury is unknown. We hypothesized that activation of PPARδ would inhibit the radiation-induced oxidative stress and proinflammatory responses in microglia. Here, we report that PPARδ activation does indeed prevent the radiation-induced increase in: i) intracellular ROS generation, ii) Cox-2 and MCP-1 protein, and iii) IL-1β and TNF-α message levels. This occurred, in part, by transrepression of NF-κB and inhibition of the PKCα/MEK1/2/ERK1/2/AP-1 pathway.

Materials and methods

Materials

The PPARδ agonist, L-165041, was purchased from Calbiochem (San Diego, CA). The MEK inhibitor, U0126, the ERK1/2 inhibitor, FR180204, and the $PKCa/\beta$ inhibitor, G06976, were purchased from EMD millipore (La Jolla, CA), Santa-Cruz Biotechnologies (Santa-Cruz, CA), and EMD millipore, respectively. All drugs were dissolved in $Me₂SO₄$ (DMSO). Goat anti-Cox-2, rabbit anti-MEK1/2, rabbit anti-p-MEK1/2, mouse anti-ERK1/2, mouse-anti-p-ERK1/2, goat anti-p-c-jun, and mouse anti-p-IκBα were purchased from Santa-Cruz Biotechnologies. Rabbit anti-p65, rabbit anti-PKCα, and rabbit anti-MCP-1 were purchased from Cell Signaling (Danvers, MA). Rabbit anti-p-PKCα was purchased from Epitomics (Burlingame, CA), and mouse anti-β-actin was purchased from Sigma-Aldrich (St. Louis, MO).

Cell culture and Irradiation

BV-2 cells, immortalized murine microglial cells, were cultured in DMEM high glucose media (Invitrogen, Carlsbad, CA) supplied with 5% fetal bovine serum (Sigma-Aldrich), 100 IU/mL penicillin, and 100 mg/mL streptomycin (Sigma-Aldrich) and were maintained at 37° C in 10% CO₂ and 90% air. Twenty-four hours prior to irradiation, the culture medium was replaced with serum-free media. Cells were irradiated with a single dose of 10 Gy using a 137Cs irradiator (J.L. Shepherd and Associates, San Fernando, CA) at a dose rate of 3.56 Gy/min. Irradiations were conducted at room temperature and control cells received sham-irradiation. Following irradiation, culture dishes were returned to the incubator and maintained at 37 \degree C in 10% CO₂ and 90% air.

Short hairpin RNA targeting

Short hairpin RNA (shRNA) were generated as described by Sui et al. [37]. The PPARδ target site is: GGACCAGAACACACGCTTCCTT. BV-2 cells were infected with ecotropic virus and infected cells were selected by puromycin selection (Sigma-Aldrich). Single-cell clones were generated, and the expression of PPARδ was evaluated by western blot analysis.

Overexpression

BV-2 cells were transfected with the pcDNA3.1.PPARδ (Invitrogen) or the pcDNA3.1 empty vector control using Lipofectamine 2000 (Invitrogen), according to the manufacturer's protocol.

Luciferase assay

Cells were plated on 24-well plates; 24 h later, cells were cotransfected with i) 0.2μ g of PPRE-ACOX (PPRE consensus sequence for the rat acyl-CoA oxidase gene, a kind gift from Dr. Thomas McIntyre, Univ. of Utah) or pGL3 control vector (Promega, Madison, WI), and ii) 0.02 μg pRL-SV40 renilla vector (a kind gift from Dr. Lee Yong Woo, Univ. of Virginia). Lipofectamine 2000 (Invitrogen) was used to transfect cells according to the manufacturer's protocol. Transfected cells were treated with vehicle control or L-165041. Twenty-four hours later, the Dual Luciferase Assay (Promega) was conducted according to the manufacturer's protocol, and the Terner Designs Reporter Microplate Luminometer (Promega) was used to measure luciferase activity. Fold change in luminescence, a measure of luciferase activity, was calculated by the relative luminescence units (RLU) of firefly/ RLU renilla luciferase.

Measurement of intracellular ROS generation

Intracellular ROS generation was measured using 2'7'-dichlorofluorescein (DCFH-DA) as previously described [38]. DCFH-DA is permeable to cell membranes; once in cells, it is cleaved by cellular esterases and becomes impermeable. When oxidized by ROS, the probe becomes fluorescent. Briefly, cells were washed with $\text{PBS} + (1 \times \text{PBS}$ supplemented with 0.14 g/L CaCl2 and 0.1 g/L of MgCl2), incubated with 10 μ M DCFH-DA (Invitrogen, CA/ Molecular Probes, Eugene, OR) for 45 min, washed with PBS+ to remove excess probe, and then incubated with 5 μM L-165041. Three hours after L-165041 treatment, cells were irradiated with a single dose of 10 Gy of $137Cs$ γ rays. One hour post-irradiation, ROS generation was measured using a FACS BDCalibur (Becton Dickinson, Bedford, MA), and BD CellQuest™ Pro 6.0 software was used to analyze the data.

RNA isolation and qRT-PCR Syber Green

RNA was harvested using Trizol reagent (Invitrogen) according to the manufacturer's protocol. DNA contamination was removed by acid-phenol chloroform extraction (pH 4.6, 125:24:1, Ambion Inc., Austin, TX). Real-time PCR amplifications were conducted in a 25 μL reaction volume containing 1 μL cDNA, 12.5 μL SYBR Green PCR Master Mix (Roche, Indianapolis, IN), 0.1 μ M upstream and downstream primers, and 10.5 μ L nuclease-free water. Real-time PCR was carried out in an ABI Prism® 7000 at 50°C for 2 min, 95°C for 2 min, and 45 cycles of 95°C for 15 min, 55°C for 30 sec, and 72°C for 30 sec. The fold changes in MCP-1, IL-1β, and TNF-α gene expression were calculated using the comparative C_t (cross threshold) method. In brief, the C_t of the housekeeping gene GAPDH was subtracted from the C_t of MCP-1, IL-1 β , or TNF- α to get Δ Ct. The ΔC_t of the sham-irradiated group was then subtracted from the ΔC_t for each of the other treatment groups to get $\Delta \Delta C_t$. Fold changes compared to the sham-irradiated group were determined by calculating $2^{-\Delta\Delta Ct}$. Data represent the mean \pm S.E.M of three independent experiments.

Immunoblotting

Total cellular protein was harvested using M-PER lysis buffer (Pierce Biotechnology, Inc., Rockford, IL) supplemented with 1 mg/mL aprotinin (Sigma-Aldrich), 1 mg/mL leupetin (Sigma-Aldrich), 10 mg/mL phenylmethylsulfonyl fluoride (PMSF), 1 mM Na_3VO_4 (Sigma-Aldrich), and 150 mM of NaCl. Lysates were centrifuged at 12,500 rpm for 10 min, and the supernatant was collected. Protein concentrations were measured using the Bradford

assay (BioRad, Hercules, CA) at absorbance 595 nm. Five to 50 μ g of protein were separated by SDS-PAGE. Protein was transferred to polyvinylidene fluoride (PVDF) membrane for 1.5–3 h at 80 V, blocked in 2.5% BSA in TBST (0.02 M Tris, 0.015 M NaCl, 0.05% Tween 20, pH 7.5), and incubated overnight with primary antibody. Membranes were washed, incubated with the appropriate HRP-conjugated secondary antibody, developed using the ECL detection system (GE Healthcare, NJ), and processed using a Kodak Processing System. Films were scanned and densitometry was conducted to quantify the signal intensity using Adobe Photoshop Elements 6.0.

Electromobility shift assay (EMSA)

Cells were lysed on ice with Buffer A (10 mM Hepes, pH 7.9, 1.5 mM $MgCl₂10$ mM KCl, 0.5 mM DTT); lysates were homogenized using a dounce homogenizer, which was followed by centrifugation at 12,000 rpm for 10 min. To extract nuclear protein, the nuclear pellets were lysed with Buffer C (5 mM Hepes, pH 7.9, 1.5 mM MgCl₂, 25% v/v glycerol, 400 mM NaCl, 1 mM EDTA 0.5 mM DTT, 0.5 mM PMSF, 2 mg/mL aprotinin, 2 mg/mL leupeptin, and 1 mM Na_3VO_4). This was followed by centrifugation at 12,000 rpm for 10 min. Protein concentrations were measured using the Bradford assay (BioRad) at absorbance 595 nm. The EMSA procedure was performed using the Promega Gel-Shift Core Assay following the manufacturer's protocol. In brief, 10μ g of nuclear protein were incubated with $2 \mu L$ Binding Buffer (Promega) for 10 min. Consensus binding sequences of NF-κB (5'- AGTTGAGGGGACTTTCCCAGGC-3' and 3'TCAACTCCCCTGAAAGGGTCCG-5') and AP-1 (5'-CGCTTGATGAGTCAGCCGGAA-3' and 3'- GCGAACTACTCAGTCGGCCTT-5') were labeled with 10μ Ci γ -P32 (GE Healthcare, Piscataway, NJ) and T4 polynucleotide kinase (Promega). The consensus sequences were then incubated with the nuclear protein for 20 min and electrophoresed on a 4% nondenaturing polyacrylamide gel. An X-ray film was then placed on top of the gel and developed overnight at −80°C. The X-ray film was processed using a Kodak Processing

Co-Immunoprecipitation

Cell lysates were collected using RIPA buffer (50mM Tris, pH 8.0, 150 mM NaCl, 1% NP-40, 0.5% deoxycholic acid, 1% SDS, 1 mg/mL aprotinin, 1 mg/mL leupetin, 10 mg/mL PMSF, and 1 mM Na_3VO_4). Lysates were incubated with 2 μg anti-p65 at 4 °C for 3 h, and immunocomplexes were captured by incubating lysates with Protein A/G PLUS agarose beads (Santa Cruz) at 4°C overnight. Agarose beads were centrifuged and washed 8 times with RIPA buffer. After centrifugation, the pellet was diluted with 80 μ 1 4× SDS-PAGE sample buffer (0.25M Tris-HCL, pH 6.8, 8% SDS, 30% Glycerol, 0.02% Bromophenol Blue, and 10% B-ME), and boiled for 5 min. The samples were then subjected to electrophoresis, and immunoblotted with an anti-PPARδ antibody.

Statistical Analysis

Each experiment was repeated a minimum of three independent times. All analyses were carried out using SAS software (SAS Inc, Cary, NC). Although the sample size within each treatment group is not large, we believe that the outcome measure distribution is normally distributed in the population. To determine statistical significance between two treatment groups, when appropriate, either a one- (when compared to sham) or two-sample t test was used. If more than two treatment groups were compared, analysis of variance (ANOVA) was used to explore the overall association. Bonferroni and Tukey's studentized range tests were used for pairwise comparisons. Levene's test was used to examine the homogeneity of variance, an assumption of variance. When this assumption was not valid, the Kruskal-Wallis test was used.

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System. Films were scanned and densitometry was performed.

Results

BV-2 cells express a functional PPARδ

To demonstrate that BV-2 cells contain a functional PPARδ and were appropriate for our studies, we performed a luciferase reporter assay. Cells were co-transfected with: i) a PPREdriven luciferase vector or a pGL3 control vector, and ii) a renilla vector. They were then incubated with the PPARδ agonist, L-165041, or vehicle control for 24 h, and luciferase activity was measured. Incubating BV-2 cells with 5 μM of L-165041 led to a 2-fold increase in luciferase activity, suggesting that these cells do express a functional PPARδ (SFig. 1). Since $5 \mu M$ of L-165041 led to a significant increase in luciferase fluorescence, we chose to use this concentration of the PPAR δ agonist for the remainder of our studies.

PPARδ activation prevents the radiation-induced increase in intracellular ROS generation and proinflammatory mediators in BV-2 cells

Radiation induces increased intracellular ROS generation in BV-2 cells [18]; we hypothesized that PPARδ activation could modulate this response. To test this hypothesis, we used the oxidation sensitive probe DCFH-DA to measure intracellular ROS generation in BV-2 cells, which were incubated with L-165041 prior to irradiation with a single dose of 10 Gy. As predicted, irradiating the cells resulted in increased ROS production 1 h postirradiation, and this response was inhibited in cells treated with L-165041 (Fig. 1A). Incubating the cells with nacetyl-cysteine, an ROS scavenger, also inhibited the radiationinduced increase in DCF fluorescence (Fig. 1A). As expected, the non-oxidizable control probe, carboxy-DCF (C369) did not show any radiation-induced difference in fluorescence (Fig. 1A).

Previous reports have demonstrated that irradiating BV-2 cells leads to an increase in Cox-2 protein and IL-1 β and TNF- α message levels [18, 19]. We hypothesized that PPAR δ activation would inhibit the radiation-induced inflammatory response in BV-2 cells. As shown in Fig. 1B (SFig. 2), L-165041 inhibited the increase in Cox-2 and MCP-1 protein levels observed 7 h post-irradiation. The increase in IL-1β and TNF-α message levels determined 24 h post-irradiation was also significantly inhibited when cells were pretreated with L-165041 (Fig. 1C).

This effect appeared to be PPARδ-dependent; L-165041 failed to inhibit the radiationinduced increase in Cox-2 in the presence of the PPARδ antagonist GSK0660 (SFig. 3) [39]. Moreover, overexpressing PPARδ by transfecting cells with the pcDNA3.1.PPARδ vector inhibited the radiation-induced increase in Cox-2 expression (Fig. 2A). In contrast, shRNAmediated knockdown of PPARδ (Fig. 2B) led to an increase in the radiation-induced Cox-2 and MCP-1 expression compared to cells infected with scrambled control shRNA. These results indicate that: i] overexpression of PPARδ prevents the radiation-induced increase in Cox-2 expression, and ii] loss of PPARδ enhances the radiation-induced increase in Cox-2 and MCP-1 expression. We also examined if PPARδ knockdown leads to an increase in IL-1β or TNF-α message levels, or intracellular ROS generation; however, we did not observe any differences in these endpoints between cells expressing scramble- or PPARδtargeted shRNA (data not shown).

PPARδ activation inhibits NF-κB activation via transrepression by physically interacting with the p65 subunit

The radiation-induced proinflammatory response in BV-2 cells is regulated, in part, by NFκB [18, 19]. Thus, we examined if PPARδ activation could modulate NF-κB activation in BV-2 cells following irradiation. As predicted, pre-incubating BV-2 cells with L-165041 prevented the radiation-induced increase in NF-κB DNA binding observed 30 min post-

irradiation (Fig 3A; SFig 4A). To investigate the mechanism by which PPARδ modulates NF-κB activation, we examined the phosphorylation of Iκ-Bα and the nuclear translocation of p65. Thirty minutes post-irradiation, we observed the predicted increases in the phosphorylation of Iκ-Bα at Ser 32 and the nuclear translocation of p65; L-165041 did not modulate these responses (Figs. 3B & 3C; SFig 4B and SFig 4C). Next, we examined if PPARδ inhibited NF-κB activation via transrepression by binding to the p65 subunit. Indeed, when we immunoprecipitated cellular lysates for p65 and then immunoblotted for PPARδ, we observed that L-165041 led to an increase in the binding of PPARδ to p65 30 min post-irradiation (Fig. 3D). Mechanistically, these data suggest that PPARδ activation prevents the DNA binding activity of NF-κB via transrepression by physically interacting with the p65 subunit.

PPARδ activation prevents AP-1 activation by inhibiting c-jun phosphorylation and upstream activation of the stress-activated kinases MEK1/2 and ERK1/2

In addition to $NF-\kappa B$ activation, activation of $AP-1$ is thought to underlie the radiationinduced increase in proinflammatory mediators in BV-2 cells [18]. We therefore investigated if L-165041 could also modulate AP-1 activation. Pre-incubating BV-2 cells with L-165041 prevented the radiation-induced increase in AP-1 DNA binding activity observed 30 min post-irradiation (Fig. 4A; SFig. 5A). Activation of AP-1, a dimeric protein, is mediated, in part, by phosphorylation of the c-jun subunit. We observed that L-165041 inhibited the phosphorylation of nuclear c-jun at Ser73 following irradiation, suggesting that PPARδ activation inhibits AP-1 activation by preventing c-jun activation (Fig. 4B; SFig. 5B).

MEK1/2 and ERK1/2 activation regulates c-jun phosphorylation and AP-1 activation in BV-2 cells [Zhiyong Deng and Weiling Zhao, unpublished data]. Interestingly, studies have demonstrated that PPARδ can modulate activation of the MEK1/2/ERK1/2 pathway [31, 40]. Thus, we hypothesized that PPARδ modulates AP-1 activation via inhibition of MEK1/2 and ERK1/2 phosphorylation. Consistent with our hypothesis, pre-incubating BV-2 cells with L-165041 reduced the radiation-induced phosphorylation of MEK1/2 at Ser218 and Ser222, and that of ERK1/2 at Thr 202 and Tyr 204 (Figs. 4C & 4D; SFigs. 5C & 5D). Furthermore, pretreating BV-2 cells with 2 μ M of the MEK1/2 inhibitor, U0126, or 2 μ M of the ERK1/2 inhibitor, FR180204, inhibited the radiation-induced increase in intracellular ROS generation and Cox-2 expression (Figs. 5A & 5B; SFigs. 6). MEK1/2 or ERK1/2 inhibition also prevented the radiation-induced increases in MCP-1, IL-1 β , and TNF- α gene expression (Figs. 6A & 6B).

PPARδ modulates MEK1/2 and ERK1/2 activation, in part, by inhibiting the radiationinduced phosphorylation and expression of PKCα

PKCα is an activator of the MEK1/2/ERK1/2 pathway [41, 42]. A radiation-induced activation of PKCα has not been previously reported in microglia. However, lipopolysaccaride (LPS), an inflammatory stimulus, has been shown to increase activation of PKCα in microglial cells [43]. Therefore, we next investigated if radiation induces PKCα activation in BV-2 cells, and further, if PPARδ activation can modulate this response. As shown in Fig. 7A (SFig. 7A), irradiating BV-2 cells with a single dose of 10 Gy led to an increase in the phosphorylation of PKCα at Thr638 and the expression of PKCα, and pretreatment with L-165041 prevented these increases.

To demonstrate that inhibition of PKCα prevents the radiation-induced activation of MEK1/2 and ERK1/2, we pre-treated BV-2 cells with 1 μM of the PKCα/β inhibitor, Go6976, and examined the phosphorylation of MEK1/2 and ERK1/2 30 min post-

irradiation. As predicted, Go6976 prevented the radiation-induced phosphorylation of MEK1/2 and ERK1/2 (Figs. 7B & 7C; SFigs. 7B & 7C).

The radiation-induced phosphorylation of MEK1/2 and ERK1/2 is modulated by ROS production in BV-2 cells [Zhiyong Deng and Weiling Zhao, unpublished data]. Our results suggest that PKCα regulates MEK1/2 and ERK1/2 phosphorylation; thus, we examined whether PKCα phosphorylation is also modulated by ROS production. Indeed, pre-treating BV-2 cells with 10 mM of NAC inhibited the radiation-induced phosphorylation and expression of PKCα (Fig. 7D; SFig. 7D). Given that L-165041 inhibited the radiationinduced increase in intracellular ROS generation (Fig. 1A), these data suggest that PPARδ activation negatively regulates the PKCα/MEK1/2/ERK1/2 pathway by preventing ROS generation following irradiation.

Discussion

A growing body of evidence suggests that PPARδ activation can regulate oxidative stress and inflammatory responses following various cellular stresses. We therefore hypothesized that PPARδ activation would prevent the radiation-induced oxidative stress and proinflammatory responses in microglia. As predicted, pretreating BV-2 cells with L-165041 inhibited the radiation-induced increase in: i) intracellular ROS generation, ii) Cox-2 and MCP-1 protein expression, and iii) IL-1β and TNF-α message levels. This occurred, in part, by negatively regulating the DNA binding of NF-κB and AP-1. PPARδ activation: i) inhibited NF-κB via transrepression by physically interacting with the p65 subunit, and ii) prevented activation of the PKCα/MEK1/2/ERK1/2/AP-1 pathway by inhibiting the radiation-induced increase in intracellular ROS generation.

The radiation-induced proinflammatory response in BV-2 cells is regulated, in part, by NFκB. There are several mechanisms by which PPARδ can modulate NF-κB, including: i) inhibiting the nuclear translocation of p65, ii) binding directly to p65, and iii) reducing acetylation of p65 [30, 44, 45]. In a rat model of reperfusion injury, the PPARδ agonist, GW0742, reduced myocardial infarct size, in part, by inhibiting nuclear translocation of p65 [44]. In cultured neonatal rat cardiomyocytes, L-165041 decreased LPS-stimulated NF-κB activation by increasing the physical interaction of PPARδ with the p65 subunit. This interaction is thought to interfere with the DNA binding of NF-κB [30]. The PPARδ agonist, GW501516, inhibited inflammation in human HaCaT keratinocytes by reducing TNF-α-induced p65 acetylation. Acetylation of p65 is regulated by the transcriptional coactivator p300; reduced acetylation of p65 decreases DNA binding and activation of NF-κB [45]. Our studies indicate that L-165041 increased the physical interaction of PPARδ to p65, suggesting that the receptor modulates NF-κB activation via transrepression.

In addition to NF-κB activation, the radiation-induced proinflammatory response in BV-2 cells is regulated, in part, by AP-1 activation. The transactivation function of the c-jun subunit of AP-1 depends on the phosphorylation of residues Ser63 and Ser73 [46]. Radiation induces phosphorylation of both residues in BV-2 cells [18]. We observed that PPARδ prevents the phosphorylation of nuclear c-jun at Ser73, and further, inhibits AP-1 DNA binding. Previous studies in BV-2 cells demonstrate that c-jun is phosphorylated by ERK1/2 [Zhiyong Deng and Weiling Zhao, unpublished data]. We observed that PPARδ activation prevents the radiation-induced activation of MEK1/2 and ERK1/2. In addition to AP-1, activation of the MEK1/2/ERK1/2 pathway has been shown to modulate NF- κ B activation. In white adipose tissue, PPARδ activation prevented LPS-induced inflammation by inhibiting ERK1/2 activation and the downstream activation of $NF- κ B [40]$. It is therefore possible that the PPARδ-mediated modulation of the MEK1/2/ERK1/2 pathway in our

model also contributes to a reduction in radiation-induced proinflammatory responses, in part, by preventing NF-κB activation.

PKC is an upstream regulator of the MEK1/2/ERK1/2 pathway. In BV-2 cells, LPSstimulated PKC activation led to ERK1/2 phosphorylation and increased Cox-2 protein levels [47]. Furthermore, in cultures of primary microglia, LPS-stimulated PKCα modulated TNF-α expression [43]. Studies have demonstrated that PPARδ can modulate PKCα. In PPARδ-null mice compared to wild-type mice, TPA induced significantly greater PKCα activity, enhancing MEK1/2 and ERK1/2 phosphorylation and Cox-2 expression [31]. Moreover, studies in platelets have demonstrated that PPARδ can physically interact with PKCα and suppress PKCα-mediated platelet activation [48]. Given the evidence that PKCα modulates both the MEK1/2/ERK1/2 pathway and inflammation, and further that PPARδ modulates PKCα activity, we examined if PKCα was an activator of MEK1/2 and ERK1/2 in our model. Indeed, we observed that the radiation-induced phosphorylation of PKCα activates the MEK1/2/ERK1/2 pathway, and PPARδ activation inhibits this response.

Our data suggest that PPARδ prevented activation of the PKCα/MEK1/2/ERK1/2/AP-1 pathway by inhibiting the radiation-induced increase in intracellular ROS generation. Of interest, MEK1/2 and ERK1/2 inhibition also prevented the radiation-induced increase in intracellular ROS generation, suggesting that the increase in intracellular ROS generation observed following irradiation reflects both direct and indirect effects, and that PPARδ activation can inhibit these responses. The antioxidant actions of PPARδ agonists have been observed in a variety of cell types [27–29]; we observed that PPARδ activation could inhibit intracellular ROS generation following irradiation of BV-2 cells. It should be noted that PPARδ can activate transcription of several antioxidant genes, including catalase, superoxide dismutase, glutathione peroxidase 1, heme oxygenase 1, and thioredoxin 1 [32, 33]. Previous studies in our lab have failed to detect changes in the expression levels of antioxidant genes in BV-2 cells; however, changes in the activity levels of antioxidant enzymes have not been examined [18]. Future studies should examine if radiation induces changes in the activity levels of antioxidant enzymes, and further, if PPARδ can modulate these changes. These studies will help elucidate the mechanisms by which PPARδ activation can modulate intracellular ROS.

Based on the findings described above, we propose a model, outlined in Fig. 8, for the role of PPARδ in modulating radiation-induced proinflammatory responses in microglia. Following irradiation, increased intracellular ROS generation leads to activation of the stress-activated kinases, PKC α , MEK1/2, and ERK1/2, and the proinflammatory transcription factors, NF-κB and AP-1. Activation of these transcription factors increases the expression of Cox-2, MCP-1, IL-1β, and TNF-α; these proinflammatory mediators contribute to the inflammatory phenotype of microglia. PPARδ activation prevents the radiation-induced proinflammatory response, in part, by negatively regulating NF-κB and AP-1. Specifically, PPARδ physically interacts with p65 and inhibits the DNA binding of NF-κB. Futhermore, PPARδ inhibits the radiation-induced increase in intracellular ROS generation, which prevents PKCα, MEK1/2, ERK1/2, and c-jun phosphorylation and the DNA binding of AP-1. Taken together, our data indicate that PPARδ activation can prevent the acute radiation-induced oxidative stress and proinflammatory responses observed in microglia within hours of *in vitro* irradiation.

It is important to note that the microglia used in our in vitro studies were grown under 21% oxygen, a concentration much higher than would occur in vivo. Thus, the affect of "physiological" oxygen concentration on the radiation-induced inflammatory response of microglial cells remains to be determined. Furthermore, although our in vitro studies examined the radiation-induced inflammatory response in microglial cells, radiation-induced

brain injury in vivo is a multicellular process. Indeed, we recognize that interpreting these data in terms of their relevance to the onset and progression of radiation-induced late effects in the brain is difficult and somewhat controversial. It is unlikely that the radiation-induced cognitive impairment observed 6 months or more after fWBI is a direct result of acute proinflammatory responses in the microglia alone. However, brain irradiation clearly leads to chronic, persistent increases in activated microglia [16, 17]; the resultant neuroinflammation has been associated with decreased neurogenesis and impaired neuronal function [15, 16, 21]. Thus, understanding the mechanisms by which radiation alters the microglia cell phenotype, and how PPARδ agonists might prevent/ameliorate these changes, offers the promise of identifying potential interventions that can be evaluated in vivo.

Moreover, since PPARδ activation has been shown to regulate inflammation in multiple cell types and promote the survival of neurons under stress conditions, it is possible that PPARδ can mitigate radiation-induced brain injury in multiple cell types [29, 49]. Additionally, animal studies have demonstrated that PPARδ agonists can cross the blood brain barrier and modulate oxidative stress and pro-inflammatory responses associated with acute and chronic CNS disorders [34–36]. Overall, our current in vitro findings indicate that PPARδ activation shows promise as a potential therapeutic strategy in the treatment and/or prevention of radiation-induced brain injury.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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List of abbreviations

WBI Whole-brain Irradiation

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Highlights

- **•** PPARδ activation prevents radiation-induced oxidative stress/inflammation
- **•** PPARδ inhibits NF-κB via transrepression by physically interacting with p65 subunit
- **•** PPARδ prevents activation of the PKCα/MEK1/2/ERK1/2/AP-1 pathway
- **•** PPARδ activation modulates radiation-induced microglial pro-inflammatory response.

Fig. 1.

PPARδ activation prevents the radiation-induced increases in intracellular ROS generation, Cox-2 and MCP-1 protein, and IL-1β and TNF-α mRNA levels. A, BV-2 cells were incubated with 10 μ M DCFH-DA or 10 μ M C369 for 45 min, the probe was washed off using PBS+, and the cells were pretreated with $5 \mu M L$ -165041, NAC, or vehicle control for 3 h. The cells were then irradiated with a single dose of 10 Gy of $137Cs \gamma$ rays or sham irradiated, and intracellular ROS generation was measured 1 h post-irradiation as described in the Materials and methods. The results are presented as arbitrary fluoresence units. B & C, BV-2 cells were pretreated with $5 \mu M$ of L-165401 and irradiated with a single dose of 10 Gy. B, Protein was harvested 7 h post-irradiation and subjected to western blot analysis for Cox-2 or MCP-1; β-actin was used as a loading control. See SFig. 2 for densitometric analysis. C, RNA was harvested 24 h post-irradiation; the expression levels of IL-1β and TNF-α were analyzed using SYBER Green Real-time PCR and normalized with GAPDH expression levels. Changes in gene expression were calculated using the $2^{\Delta\Delta Ct}$ methods (see Materials and methods). Mean \pm S.E.M; *, p 0.05 vs. sham-irradiated, #, p 0.05 vs. 10 Gy; $n=3$.

Fig. 2.

Overexpression or knockdown of PPARδ modulates Cox-2 and MCP-1 expression in BV-2 cells. A, BV-2 cells were transfected with 24 μg of the pcDNA3.1PPARδ vector or 24 μg of the empty pcDNA3.1 vector using Lipofectamine 2000 according to the manufacturer's protocol. Twenty-four h post-transfection, cells were serum starved for 24 h, treated for 3 h with 5 μ M L-165041 or vehicle control, irradiated with a single dose of 10 Gy, and protein was harvested 7 h post-irradiation. Protein was subjected to western blot analysis for Cox-2; β-actin was used as a loading control. B, BV-2 cells were infected with scramble control or shRNA targeting PPARδ. Single knockdown clones were selected, irradiated with a single dose of 10 Gy, and protein was harvested 7 h post-irradiation. Protein was subjected to western blot analysis for Cox-2 or MCP-1. β-actin was used as a loading control.

Fig. 3.

PPARδ activation prevents radiation-induced NF-κB activation by physical interaction with the p65 subunit. BV-2 cells were pretreated with 5μ M of L-165401 or vehicle control and irradiated with a single dose of 10 Gy. A, Nuclear protein was collected 30 min postirradiation. Gel-Shift analysis was carried out by incubating 10μ g of nuclear protein with γ -ATP P^{32} end- labeled NF- κ B consensus oligo (see Materials and methods). The samples were run on a 4% non-denaturing acrylamide gel, stained with 7% Acetic acid, and exposed to X-ray film. B, Protein was harvested 30 min post-irradiation; whole cell lysates were subjected to western blot analysis for p-IκBα. C, Nuclear protein was harvested 30 min post-irradiation; nuclear protein was subjected to western blot analysis for nuclear p65. B & C, β-actin was used as a loading control. A–C, See SFig. 4 for densitometric analysis. D, Co-immunoprecipitation was carried out using anti-p65 and immunoblotting for PPARδ (see Materials and methods).

Fig. 4.

PPARδ activation prevents radiation-induced AP-1 activation and upstream activation of cjun, MEK1/2, and ERK1/2. BV-2 cells were pretreated with $5 \mu M$ of L-165041 or vehicle control and irradiated with a single dose of 10 Gy. A, Nuclear protein was collected 30 min post-irradiation. Gel-Shift analysis was carried out by incubating 10μ g of nuclear protein with γ-ATP P^{32} end- labeled AP-1 consensus oligo (see Materials and methods). The samples were run on a 4% non-denaturing acrylamide gel, stained with 7% Acetic acid, and exposed to X-ray film. B, Nuclear protein was collected 30 min post-irradiation; nuclear protein was subjected to western blot analysis for p-c-jun; β-actin was used as a loading control. C & D, Protein was harvested 30 min post-irradiation and subjected to western blot analysis for C, p-MEK1/2 and total-MEK1/2 and D, p-ERK1/2 and total-ERK1/2; β-actin was used as a loading control. A–D, See SFig. 5 for densitometric analysis.

Fig 5.

The MEK1/2 inhibitor, U0126, or the ERK1/2 inhibitor, FR180204, prevent the radiationinduced increase in intracellular ROS generation and Cox-2 expression. A, BV-2 cells were incubated with 10 μ M DCFH-DA for 45 min, the probe was washed off using PBS+, and the cells were pretreated with 2 μM U0126, 2 μM FR180204, or vehicle control for 3 h. The cells were then irradiated with a single dose of 10 Gy of $137Cs \gamma$ rays or sham irradiated, and intracellular ROS generation was measured 1 h post-irradiation. B, BV-2 cells were pretreated with 2 μM U0126 or 2 μM FR180204, or vehicle control for 1 h prior to treatment with 5 μM L-165041 or vehicle control. Three hours post-L-165041 treatment, cells were irradiated with a single dose of 10 Gy and whole cell lysates were collected. Protein was subjected to western blot analysis for Cox-2. β-actin was used as a loading control. See SFig. 6 for densitometric analysis.

Fig. 6.

MEK1/2 or ERK1/2 inhibition prevents the radiation-induced increase in MCP-1, IL-1β, and TNF-α gene expression. BV-2 cells were pretreated with A, 2 μM U0126, B, 2 μM FR180204, or vehicle control for 3 h. Three h post-treatment, cells were irradiated with a single dose of 10 Gy and RNA was harvested 24 h post-irradiation. The expression levels of MCP-1, IL-1β, and TNF-α were analyzed using SYBER Green Real-time PCR and normalized with GAPDH expression levels. Changes in gene expression were calculated using the $2^{-\Delta\Delta Ct}$ methods (see Materials and methods). Mean \pm S.E.M; *, p 0.05 vs. shamirradiated, $#$, p 0.05 vs. 10 Gy; n=3.

Fig. 7.

PPARδ modulates MEK1/2 and ERK1/2 activation, in part, by inhibiting upstream activation of PKCa. BV-2 cells were pretreated with A, 5 μ M L-165041, B & C, 1 μ M Go6976, D, 10 mM NAC, or vehicle control 3 h prior to a single dose of 10 Gy. Protein was harvested 30 min post-irradiation and subjected to western blot analysis for A & D, p-PKCα and total-PKCα, B, p-MEK1/2 and total-MEK1/2, or C, p-ERK1/2 and total-ERK1/2. A–D, β-actin was used as a loading control. A–D, See SFig. 7 for densitometric analysis.

Fig. 8.

Proposed model outlining the role of PPAR δ in the modulation of the radiation-induced proinflammatory response in BV-2 cells. Irradiation of BV-2 cells leads to an increase in intracellular ROS generation. This increases activation of the stress-activated kinases, PKCα, MEK1/2, and ERK1/2, and the proinflammatory transcription factors, NF-κB and AP-1. Activation of NF-κB and AP-1 enhance the expression of Cox-2, MCP-1, IL-1β, and TNF-α. PPARδ activation prevents radiation-induced neuroinflammation, in part, by transrepression of NF-κB by physical interaction with the p65 subunit. Additionally, PPARδ inhibits intracellular ROS generation, which prevents PKCα, MEK, and ERK1/2 phosphorylation and downstream activation of AP-1.