

**ORIGINAL RESEARCH** 

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# A Lipid-Based Oral Supplement Protects Skin Cells in Culture from Ultraviolet Light and Activates Antioxidant and Anti-Inflammatory Mechanisms

Steven R. Hall, PhD<sup>1,6</sup>, Anna-Jean Reid, MSc<sup>6</sup>, Jasmine Eng<sup>1</sup>, Brendan T. McKeown, MSc<sup>2,5</sup>, Marc St-Onge, BSc<sup>6</sup>, Kerry B. Goralski, PhD<sup>1-5\*</sup>

## **ABSTRACT**

Overexposure to ultraviolet (UV) light is associated with multiple health risks, from sunburn and prematurely aging skin to the development of skin cancers. The ingestion of photoprotective natural compounds through diet or supplementation is one method to increase the skin's UV-resistance. This study's primary objective was to determine the cellular photoprotective properties of an ingestible skincare supplement (trade name "Anti-Aging Formula" [AAF]) and compare them to its constituent active ingredients: fish oil-derived omega-3s eicosapentaenoic acid (EPA) and docosahexaenoic acid (DHA), borage-derived omega-6 gamma-linolenic acid (GLA), paprika- and marigold-derived carotenoids, zeaxanthin and lutein, respectively, and vitamin D3. AAF, but not the separate individual ingredients, significantly increased the viability of primary human dermal fibroblasts after UVA exposure compared to the vehicle control. AAF and EPA/DHA-containing fish oil demonstrated similar UVB photoprotective properties whereas GLA, the carotenoids, and vitamin D3 had no significant effect. The second objective was to explore possible mechanisms of action of AAF's photoprotective effects. AAF-treatment increased cellular antioxidant activity and the expression of genes in the glutathione and peroxiredoxin (PRDX)/thioredoxin (TXN) antioxidant pathways, suggesting an antioxidant mechanism of action. It also diminished cellular arachidonic acid (AA) levels and decreased the expression of the downstream pro-inflammatory prostaglandin-endoperoxide synthase 2 (PTGS2) gene, suggesting an anti-inflammatory mechanism of action. In conclusion, AAF is UVA/B photoprotective when applied directly to primary human dermal fibroblasts. In addition, its photoprotective effects are mainly due to its EPA/DHA components and may relate to its cellular antioxidant effects and inhibition of the AA/PTGS2 inflammatory pathway.

**KEYWORDS:** Antioxidants; borage oil; cell culture; dietary lipids; fish oil; inflammation; lutein; omega-3 and -6 fatty acids; photoprotection; zeaxanthin

## Introduction

The cells that make up our skin protect us from our environments' stressors, such as pathogens, pollution, and UV light. Overexposure to UV radiation causes a

number of premature aging effects, including skin wrinkling, loss of elasticity, or age spots, and is considered a "complete carcinogen" due to its ability to initiate and promote the growth of cancerous tumours by damaging

<sup>&</sup>lt;sup>1</sup>College of Pharmacy, Faculty of Health, Dalhousie University, Halifax, Nova Scotia, Canada B3H 4R2

<sup>&</sup>lt;sup>2</sup>Department of Pharmacology, Faculty of Medicine, Dalhousie University, Halifax, Nova Scotia, Canada B3H 4R2

<sup>&</sup>lt;sup>3</sup>Department of Pediatrics, Faculty of Medicine, Dalhousie University, Halifax, Nova Scotia, Canada B3H 4R2

<sup>&</sup>lt;sup>4</sup>Division of Hematology/Oncology, IWK Health Centre, Halifax, Nova Scotia, Canada B3K 6R8

<sup>&</sup>lt;sup>5</sup>Beatrice Hunter Cancer Research Institute, Halifax, Nova Scotia, Canada B3H 4R2

<sup>&</sup>lt;sup>6</sup>Bend Beauty, Inc., Halifax, Nova Scotia, Canada B3J 0C7

<sup>\*</sup> Corresponding Author: kerry.goralski@dal.ca



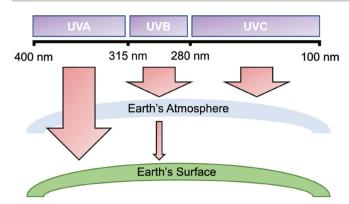


Figure 1. Types of UV radiation. The spectrum of UV radiation is divided into UVA (315-400 nm) and UVB (280-315 nm), which reach the earth's surface, and UVC (100–280 nm), which is blocked by the earth's atmosphere. Figure adapted from D'Orazio, et al. [1].

and mutating skin cell DNA [1-4]. Fibroblasts, found in the dermis layer, are considered the key drivers of the ageing process as they are quiescent, reside in the skin for years, and accumulate macromolecular damage over time [5-7].

The UV radiation that reaches the Earth's surface (and our skin) is comprised of predominantly UVA (90-95%) with a lesser amount of UVB (5-10%), both of which can induce photoaging and photocarcinogenesis (Figure 1) [1, 3]. The damaging effects of UVB occur predominantly in the epidermal skin layer [1] and result from direct damage and lesions to DNA [8] while the mutagenic and carcinogenic effects of UVA, which penetrates deeply into the dermal layer [1], primarily result from the generation of reactive oxygen species (ROS) [9] which react with and damage DNA, proteins, and lipids [10].

While the best way to reduce the damaging effects of UV light is to avoid excessive exposure by staying indoors, using sunscreen, or wearing sun-protective clothing [11], an additional method is through dietary consumption or supplementation of compounds called photoprotectors that systemically increase the skin's resistance to UV radiation. Certain photoprotectant molecules can enhance DNA repair, reduce UV-induced immunosuppression, ROS activity, and even the incidence of skin cancers [12].

AAF contains a mixture of the omega-3 long chain polyunsaturated fatty acids (LCPUFAs) EPA and DHA, the omega-6 LCPUFA GLA, and the carotenoids lutein and zeaxanthin, which individually display anti-inflammatory or anti-oxidant properties [13–16]. Ingestion of EPA and DHA has also been shown to clinically reduce sunburn, skin photosensitivity, photoageing, and photocarcinogenesis [13, 17] and the ingestion of the carotenoids lutein and zeaxanthin clinically reduce ocular

[18] and dermal phototoxicity [16, 19]. In a clinical trial, daily ingestion of the commercial skincare supplement AAF was photoprotective based on an average 39% and 84% increase in minimal erythema dose (MED; the smallest amount of UV radiation that produces skin redness) after 4 and 8 weeks, respectively. Additionally, 100% of the trial's 28 participants had higher MED values after 8-weeks of supplementation versus their baseline values [20].

In an effort to better understand the photoprotective activity of AAF at the cellular level, we aimed to: develop methodology to quantify UVA/B photoprotection using human dermal fibroblasts, compare the photoprotective effects of AAF versus its individual components, and develop insight into AAF's cellular photoprotective mechanisms of action.

## **Materials & Methods**

#### **Materials**

AAF was obtained from Anti-Aging Formula capsules (Bend Beauty, Inc., Halifax, NS, Canada). AAF contains 0.91 M EPA and 0.28 M DHA derived from fish oil (FO), 0.11 M GLA from borage seed oil (BSO), 2.3 mM lutein from marigold flower extract (MFE; Tagetes erecta), 1.2 mM zeaxanthin from paprika extract (PE; Capsicum annuum), and 17 µM vitamin (Vit) D3, with the remaining non-medicinal components being beeswax, olive oil, sunflower lecithin, natural mint flavouring, safflower oil, tocopherols, ascorbyl palmitate, and medium chain triglycerides, encapsulated in bovine gelatin capsules. Molecular biology grade water, fibroblast medium (Medium 106), low serum growth supplement (LSGS), fetal bovine serum (FBS), penicillin and streptomycin, H<sub>2</sub>O<sub>2</sub>, 5-(and 6-)chloromethyl-2'7'-dichlorodihydrofluorescein diacetate (CM-DCFH<sub>2</sub>-DA), trypsin/EDTA, trypsin neutralizer, Alexa Fluor 488 Phalloidin, Super Script II Reverse Transcriptase, and a handheld e-series UVA<sub>365</sub>/UVB<sub>312</sub> lamp were purchased from Thermo Fisher Scientific (Burlington, Ontario, Canada). A radiometer with UVA/B sensors was purchased from Solar Light Company (Glenside, Pennsylvania, USA). SsoAdvanced Universal SYBR Green Supermix and Aurum Total RNA Mini Kit was purchased from Bio-Rad (Mississauga, Ontario, Canada). Dako Fluorescent Mounting Medium was purchased from Agilent (Santa Clara, California, USA). All other chemicals were purchased from Sigma Aldrich (Oakville, Ontario, Canada).

## Cells

Primary human dermal fibroblasts (C0135C; Cascade Biologics) were cultured in Medium 106 supplemented with LSGS and 100 IU/mL penicillin, 250 μg/mL



streptomycin (complete fibroblast medium). Cells were split weekly and medium changed twice a week. Cells were maintained in a humidified, 95% air/5% CO<sub>2</sub> atmosphere at 37°C (standard conditions) for a maximum of three weeks (six passages), after which they began to senesce.

#### Dissolution of AAF

Carotenoids, such as lutein and zeaxanthin, are known to be difficult to use in cell culture studies due to their poor solubility in aqueous solutions [21]. With percentages (v/v) relative to the total solution volume, 10% AAF (or 10% ddH<sub>2</sub>O for vehicle control) was dissolved in 10% tetrahydrofuran (THF) [22] supplemented with 0.00125% butylated hydroxytoluene (BHT) [23], 75% dimethyl sulfoxide (DMSO), and 5% FBS [23–25].

## **AAF Fibroblast Dose Determination**

Thiazolyl blue methyltetrazolium bromide (MTT) cell viability assays were completed in AAF-treated fibroblasts (0.0025–0.04%), as previously described [26] with the minor alteration that the MTT reaction was 4 hours. The maximum tolerated concentration (MTC) is defined as the highest AAF concentration that did not significantly reduce cell viability; 50% of MTC (MTC<sub>1</sub>) was then used for all subsequent experiments [27].

## **Photoprotection Trials**

Fibroblasts were treated with the MTC<sub>14</sub> of AAF (0.005%), equivalent concentrations of individual active ingredients (EPA-/DHA-containing FO, GLAcontaining BSO, lutein-containing MFE, zeaxanthin-containing PE, or Vit D3), or vehicle for up to 13 days at standard conditions.

## MTT assays

Zero-, 5-, and 12-day treated cells were seeded (10,000 cells/well, 96-well plate), left to adhere overnight in 0.005% AAF treatment-, the AAF vehicle, or no-treatment containing medium, re-treated the following day and again left overnight until the target treatment times were reached. The medium was aspirated and replaced with 100–200 mL of treatment-free complete fibroblast medium (UVA assays) or 50 µL phosphate buffered saline (PBS; UVB assays). The cells were exposed to UVA (216-324 J/cm<sup>2</sup>, ~12-18 hours), UVB (31.25-24,000  $mJ/cm^2$ , ~0.2–30 minutes), or not exposed (baseline) at standard conditions. The UV-exposed medium was replaced with treatment-containing complete fibroblast medium, and the cells were left to recover overnight. MTT assays were then completed as described above. The percent-cell viability was calculated by normalizing to the baseline values.

# Fluorescent microscopy assays

Thirteen-day AAF-treated fibroblasts seeded on  $25 \times 75 \times 1.0$  mm coverslips inside 6-well plates [28] (400,000 cells/well) were left to adhere overnight in 0.005% AAF treatment- or AAF vehicle-containing medium. The medium was then replaced with 1 mL treatment-free complete fibroblast medium (UVA assays) or PBS (UVB assays), and plates were exposed to 100 J/cm<sup>2</sup> UVA, 1 J/cm<sup>2</sup> UVB, or No UV (baseline). As per published methods, the cells were then treated with 5 mg/mL of the cell-death indicator propidium iodide (PI), fixed in 4% Paraformaldehyde, treated with 1 U methanolic Alexa Fluor 488 phalloidin (F-actin stain), and 1 µg/mL Hoechst 33258 (membranepermeable nucleic dye) [29, 30]. Cell-adhered coverslips were mounted on microscope slides using Dako Fluorescent Mounting Medium and images taken using a Zeiss LSM 710 confocal microscope ( $\mathrm{Ex}_{\mathrm{405}}/\mathrm{Em}_{\mathrm{405-485}}$ for Hoechst 33258,  $Ex_{488}/Em_{495-550}$  for phalloidin,  $Ex_{561}/Em_{495-550}$ Em<sub>565-620</sub> for PI). Percent of dead versus total cells was calculated using Equation 1.

% of dead versus total cells

$$= \left(\frac{\text{# of PI stained nuclei}}{\text{# of Hoechst } 33258\text{-stained nuclei}}\right) \times 100\% \quad (1)$$

## **Antioxidant Trials**

Fibroblasts were treated with AAF or AAF vehicle for up to 14 days and seeded the same way as described in Photoprotection Trials, MTT assays. After the target treatment times were reached the media was aspirated and replaced with complete fibroblast medium containing H<sub>2</sub>O<sub>2</sub> (11.1-2,700 µM) or equivalent ddH<sub>2</sub>O vehicle for 48 hours. MTT assays were completed, and half-maximal inhibitory concentrations (IC<sub>50</sub> values) calculated as measures of H<sub>2</sub>O<sub>2</sub> cytotoxic potency [26].

To quantify AAF's effects on intracellular ROS activity, fibroblasts were seeded at 10,000 cells/well in black-sided clear-bottomed 96-well plates, left overnight at standard conditions, then treated with 7.5 μM CM-DCFH<sub>2</sub>-DA for 1 h in complete fibroblast medium. This solution was aspirated, the wells washed with PBS, and the cells treated with 900 mM H<sub>2</sub>O<sub>2</sub> or ddH<sub>2</sub>O vehicle in complete fibroblast media for 1 hour, after which fluorescence was read (Ex<sub>485</sub>, Em<sub>528</sub>) as a measure of intracellular ROS activity [31]. To determine if there was a direct chemical interaction between AAF ingredients and H<sub>2</sub>O<sub>2</sub> the assay was repeated in the absence of fibroblasts where AAF (0.005%) or AAF vehicle in complete fibroblast medium was mixed with 900 mM H<sub>2</sub>O<sub>2</sub> or ddH<sub>2</sub>O and 3.75 μM CM-DCFH<sub>2</sub>-DA for 1 hour. The fold-increases in ROS activity were calculated as follows:



Fold increase = 
$$\frac{F_{\text{avg,treatment}}}{F_{\text{avg,vehicle}}}$$
 (2)

Where  $F_{\rm avg,treatment}$  is the average fluorescence in the  ${\rm H_2O_2}$ -treated wells and  $F_{\rm avg,vehicle}$  is that of the ddH<sub>2</sub>Otreated wells [31].

## Cellular Lipid Profile Analysis

Fibroblasts grown for 14 days in 0.005% AAF or vehicle were stored in 12 mL 2:1 chloroform:methanol with 0.01 mg internal standard tricosanoic acid (23:0). Cells were lysed by sonication and lipids extracted using a modified Folch method [32]. Extracted lipids were converted to methyl esters using 0.5 N H<sub>2</sub>SO<sub>4</sub> in methanol (100 °C, 1 hour) [33]. Fatty acid analysis was performed on recovered samples (0.10 mg/mL) using splitless injection (250 °C injector temperature) on a gas chromatograph (Bruker) with DB-23 column (Agilent Technologies) and flame ionization detector (GCFID). Temperature program used: initial temperature (60 °C, 0.5 minutes), increased to 150 °C at 45 °C/min, temperature held for 2 minutes, then increased (5.1 °C/min) to a final temperature of 220 °C and held for 5.77 minutes; 24-minute total run time. FID was set to 270 °C. Samples analyzed in triplicate. Fatty acid profiles reported as mass percent of total fatty acid identified.

# RNA Isolation, Reverse Transcription, and Quantitative Polymerase Chain Reaction (qPCR)

Quantitative PCR was performed on total RNA isolated from 14-day AAF- (0.005%) or vehicle-treated fibroblasts 6 hours after UVA- (100 J/cm<sup>2</sup>) or UVB-(1.5 J/cm<sup>2</sup>) exposure, or baseline-controls, as previously described [29]. Table 1 contains the gene-specific primers used.

## Statistical Analysis

All data are presented as mean  $\pm$  standard error of the mean (SEM) of at least three independent trials.

Unpaired t-tests were performed for dual comparisons. One-way or two-way analysis of variances (ANOVAs) were performed for multiple comparisons with one or two independent variables, respectively, and followed by Dunnett's or Tukey's multiple comparisons tests, respectively. A difference was considered significant if  $P \le 0.05$ .

## Results

# **AAF-Treatment Protects Human Dermal** Fibroblasts from UVA/B-induced Cell Death

Treatment with AAF (0.02%) reduced fibroblast viability to 52%. Therefore, AAF's MTC was 0.01% and the concentration used in all subsequent trials, the MTC<sub>1/2</sub>, was 0.005% (Figure 2A). UVA exposure (216 J/cm<sup>2</sup>) reduced the viability of the vehicle-treated fibroblasts to 19%, which increased to 55%, 80%, and 66% when the fibroblasts were pre-treated with 0.005% AAF for 1-, 7-, or 14-days, respectively (Figure 2B). UVB's IC<sub>50</sub> quantified via MTT cell viability assays was 1.7-fold higher in 14-day AAF- versus vehicle-treated fibroblasts (Figure 2C) versus vehicle-treated cells, fibroblasts pre-treated with AAF were 31% and 11% more viable after UVA or UVB exposures, respectively (Figures 2D–E). None of the individual components significantly increased cell viability after UVA exposure; only FO significantly increased cell viability after UVB exposure, by 10%.

# AAF Helps Preserve Fibroblast Morphology Following UVA/B Exposure

Fibroblasts treated with vehicle or AAF for 14 days had intact, elongated actin fibers and no PI-stained nuclei (Figure 3A, left). In comparison, the vehicle-treated cells exposed to UVA or UVB had a diffuse F-actin staining pattern and lack of elongated actin fibers, reduced fibroblast density and size, and brightly red PI-stained nuclei indicating cell death (Figure 3A, top middle and right). The AAF- and UVA-treated cells demonstrated

Table	<b>1.</b> PCR pri	mers used to	determine th	e expression (	ot relevan	t genes ir	n primary	human d	lermal fibroblasts.
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Gene	PCR forward primers (5'-3')	PCR reverse primers (5'-3')
PTGS1	GCACAGGAGCCTGCACTC	GTCACACTGGTAGCGGTCAA
PTGS2	CTGATGATTGCCCGACTCCC	CGCAGTTTACGCTGTCTAGC
GCLC	GGAAGTGGATGTGGACACCAGA	GCTTGTAGTCAGGATGGTTTGCG
GSS	GGAGCCTCTTGCAGGATAAAC	GAGACGAGCGGTAAAGTC
GSR	GAGATGGCAGGGATCCTGTCAGC	ATTCTGGAATTCGTCTACGATGATATGACC
PRDX1	TTTGGTATCAGACCCGAAGC	TCCCCATGTTTGTCAGTGAA
TXN	GGTGAAGCAGATCGAGAGCA	CCACGTGGCTGAGAAGTCAA
TXNRD1	CCACTGGTGAAAGACCACGTT	AGGAGAAAAGATCATCACTGC
GAPDH	GAGTCAACGGATTTGGTCGT	TTGATTTTGGAGGGATCTCG
ACTB	GGACTTCGAGCAAGAGATGG	AGCACTGTGTTGGCGTACAG

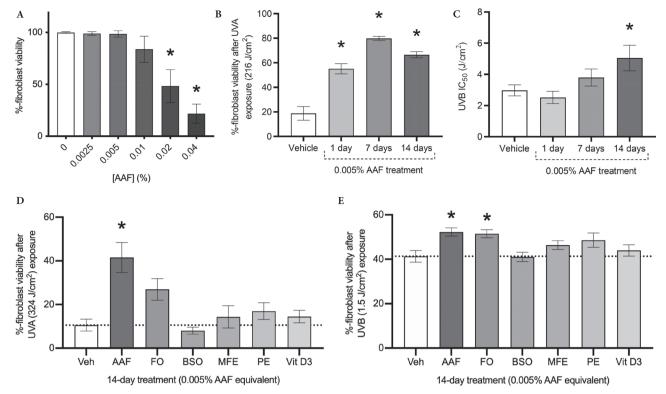


Figure 2. AAF is photoprotective in human primary dermal fibroblasts. (A) Primary human dermal fibroblasts were treated with varying concentrations of AAF to establish the MTC. The fibroblasts were then treated with AAF at the determined MTC<sub>4.7</sub> (0.005%) or vehicle for 1-, 7-, or 14-days followed by (B) UVA exposure (216 |/cm<sup>2</sup>) or (C) UVB exposure (31.25–24,000 mJ/cm<sup>2</sup>) followed by MTT cell viability-quantifying assays to evaluate its cellular photoprotective effects. Fibroblasts were treated with the AAF vehicle (Veh), AAF, or the formula's individual active components alone: EPA- and DHA-containing fish oil (FO), GLA-containing borage seed oil (BSO), lutein-containing marigold flower extract (MFE), zeaxanthin-containing paprika extract (PE), or vitamin D3 (Vit D3) for 14-days followed by (D) UVA-exposure (324 |/cm<sup>2</sup>) or (E) UVB-exposure (1.5 |/cm<sup>2</sup>). \*Signifies that the value is significantly different from that of the vehicle-treatment control in that figure, as calculated using one-way ANOVAs followed by Dunnett's multiple comparisons test (P < 0.05).

similar morphology to the No UV (baseline) control cells, albeit with a small reduction in cell density, while the AAF- and UVB-treated cells had decreased cell density, more diffuse F-actin staining, and some faint PI-staining (Figure 3A, bottom middle and lower right). Staining with the cell membrane-permeable Hoechst 33258 (blue) provided a total cell count (Figure 3B). Using Equation (1), the percentage of dead versus total fibroblasts was 98% for vehicle-treated versus 2% for AAF-treated cells after UVA exposure, and 87% for vehicle-treated versus 6% for AAF-treated cells after UVB exposure (Figure 3C).

# **AAF-treatment Protects Fibroblasts** from the ROS, H<sub>2</sub>O<sub>3</sub>

The IC<sub>50</sub> value for H<sub>2</sub>O<sub>2</sub> increased from 221 µM in vehicle-treated fibroblasts to 435, 537, and 558 μM after 1-, 7-, and 14-day AAF-treatments (Figure 4A). H<sub>2</sub>O<sub>2</sub>-treatment increased intracellular ROS activity by 8.8-fold relative to the H<sub>2</sub>O<sub>2</sub>-vehicle (ddH<sub>2</sub>O) in the AAF vehicle pre-treated cells (Figure 4B). This fold-increase was reduced to 4.5-, 4.4-, and 4.3-fold in the 1-, 7-, and 14-day AAF-treated fibroblasts. H<sub>2</sub>O<sub>2</sub> directly oxidized the ROS-detecting probe, CM-DCFH<sub>3</sub>-DA, in cell-free fibroblast medium, increasing the fluorescent reading by 55-fold in both the presence and absence of AAF (Figure 4C).

## AAF-treatment Alters the Fibroblasts' Lipid **Profile**

Representative GCFID chromatograms (Figures 5A-B) and analysis of triplicate fatty acid mass percent data (Figures 5C-F) show that treating the fibroblasts with AAF for 14-days enriched GLA, EPA, and DHA by 3.1-, 5.8-, and 1.2-fold, and diminished AA by 1.4-fold to 0.7× of what was quantified in the vehicle-treated cells, without significantly changing total fatty acid content (Figure 5G).

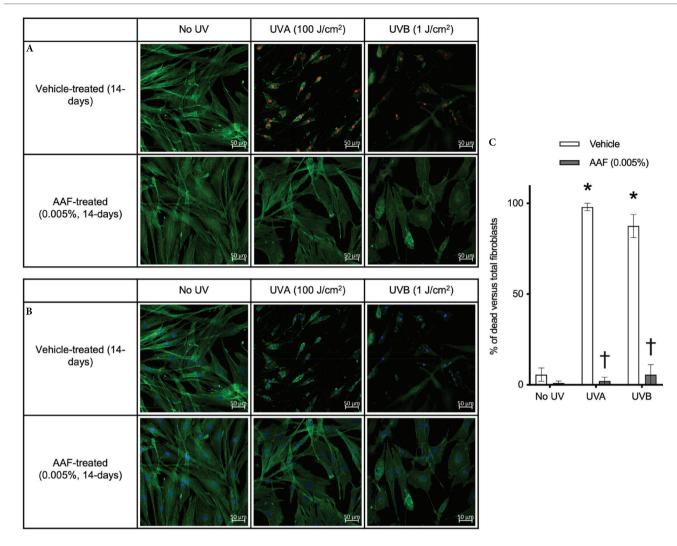


Figure 3. AAF preserves cellular morphology and prevents cell death following UVA and UVB exposure. Human primary dermal fibroblasts were treated with AAF at its MTC<sub>1/2</sub> (0.005%) for 14 days followed by UVA (100 J/cm²) or UVB (1 J/cm²) exposure. (A) Confocal fluorescent microscopy was used to visualize the effect of AAF treatment and UV-exposure on fibroblast actin cytoskeleton (phalloidin-stained F-actin [green]) and on fibroblast death (PI-stained nuclei [red]). (B) The same fibroblasts were also stained with cell membrane-permeable blue Hoechst 33258 nuclei stain to provide total cell count. (C) The percent of dead versus total cells was quantified using Equation 1. \*Signifies the value is significantly higher than the vehicle-treated baseline (No UV) control cells and ⁺signifies value is significantly lower than that of the vehicle-treated cells in the same UV-treatment group, as measured using a two-way ANOVA followed by Tukey's multiple comparisons test (P < 0.05).

# **AAF-treatment Alters the Expression of Genes Involved in Inflammation and Redox Control**

We evaluated three glutathione antioxidant pathway genes, glutamate-cysteine ligase catalytic subunit (GCLC), glutathione synthetase (GSS) and glutathione reductase (GSR), three PRDX/TXN antioxidant pathway genes, peroxiredoxin 1 (PRDX1), thioredoxin (TXN), and thioredoxin reductase (TXNRD1), and two AA/PTGS inflammatory signaling pathway genes, prostaglandin-endoperoxide synthases 1 and 2 (PTGS1/2; also known as cyclooxygenases [COX] 1/2).

Fibroblasts treated with AAF for 14 days had increased GCLC expression versus vehicle-treatment in the No UV baseline control (Figure 6A). UVA treatment increased GCLC expression compared to baseline and the increase was significantly greater in the AAF- versus vehicle-treated cells. GSS and GSR were not significantly affected by AAF treatment alone or by UVA treatment with or without AAF (Figure 6B & C). In comparison, UVB treatment decreased the expression of all three glutathione antioxidant genes relative to baseline in the vehicle-treated cells. These UVB-mediated decreases in gene expression were not reversed by AAF treatment.

AAF treatment increased the expression of *TXNRD1* but not PRDX1 or TXN compared to vehicle treatment in the baseline control cells (Figure 6D, E & F).

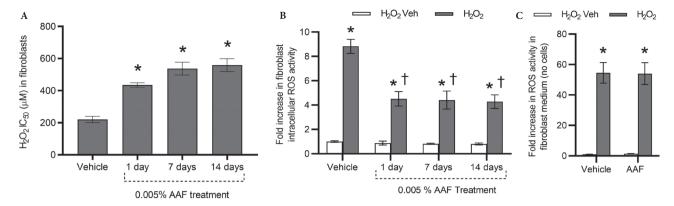


Figure 4. AAF reduces cytotoxicity and intracellular ROS-activity induced by H<sub>2</sub>O<sub>2</sub>. Primary human dermal fibroblasts were treated with AAF (0.005%) for 1-, 7-, or 14-days or AAF vehicle (Vehicle) for 14-days. This was followed by (A) treatment with various doses of the H<sub>2</sub>O<sub>2</sub> or ddH<sub>2</sub>O (H<sub>2</sub>O<sub>2</sub> Veh) after which MTT assays were performed and H<sub>2</sub>O<sub>2</sub>'s IC<sub>50</sub> values determined, or (B) treatment with the ROS-detecting fluorescent probe, CM-DCFH<sub>2</sub>-DA (7.5 mM, 1 h), after which cells were rinsed with PBS and then exposed to a single dose of H<sub>2</sub>O<sub>2</sub> (900 mM,  $\bar{1}$  h). Fluorescence was read (Ex<sub>485</sub>, Em<sub>528</sub>) as a measure of intracellular ROS activity using Equation 2. (C) In complete fibroblast medium with no cells,  $H_2O_2$  (900 mM, 1 h) or  $H_2O_2$  Veh was mixed with 3.75  $\mu$ M CM-DCFH<sub>3</sub>-DA and 0.005% AAF or AAF-vehicle (Vehicle), total ROS quantified, and fold-increase calculated *via* Equation 2. \*Signifies the value is significantly different from the vehicle control as measured using one-way ANOVAs followed by Dunnett's multiple comparisons test (A), or the H<sub>2</sub>O<sub>2</sub>-Veh within each treatment group (B & C) as measured using a two-way ANOVA followed by Tukey's multiple comparisons test. †Signifies the value is significantly different from the AAF-Vehicle/H,O,-treated cells as measured using a two-way ANOVA followed by Tukey's multiple comparisons test. P < 0.05 for all analyses.

In UVA-exposed cells, with or without AAF treatment, there was a significant increase in PRDX1, TXN, and TXNRD1 expression relative to the respective baseline controls. In UVB-treated cells, with or without AAF treatment, there was a significant increase in PRDX1 and TXN and a significant decrease in TXNRD1 relative to the respective baseline controls.

AAF treatment did not affect PTGS1 or PTGS2 expression in the baseline control cells. UVA-treatment dramatically increased (130-fold) PTGS2 but not PTGS1 expression in the vehicle-treated cells (Figure 6G & H). The UVA-mediated induction of *PTGS2* was reversed by AAF treatment. UVB treatment, with or without AAF, reduced *PTGS1* but not *PTGS2* expression.

## Discussion

Primary human dermal fibroblasts were chosen for this study because they are an accepted cell model for photoprotection trials [36–38] and a powerful preclinical tool for investigating changes in normal skin physiology [39]. Building on our previous demonstration of skin photoprotection in humans [20] we have now determined that AAF exerts UVA and UVB photoprotective effects when applied directly to primary human dermal fibroblast cells in culture. Since AAF and not the individual ingredients provided significant UVA photoprotection it suggests that the combined effects of the individual

photoprotective components (EPA & DHA, lutein, and zeaxanthin [13, 18, 40]) are required to mediate the full UVA photoprotective effect of AAF. For both UVA and UVB photoprotection, EPA/DHA containing FO appears to be AAF's primary photoprotective ingredient within this cell culture model.

In the fluorescent microscopy experiments, the lack of PI-stained nuclei in the AAF-treated/UVAexposed cells provide additional evidence of AAF's UVA-photoprotective effects; however, as these cells appeared less dense than the baseline control cells, it did not appear to completely prevent photodamage at the 100 J/cm<sup>2</sup> UVA dose used, which is consistent with the results of the MTT assays. AAF-treated/UVBexposed cells also displayed less PI-staining than the vehicle-treated/UVB-exposed cells, further supporting that AAF was also UVB photoprotective; however, since the AAF-treated/UVB-exposed fibroblasts appeared more rounded and less fibrous than those exposed to UVA, this suggests that the UVB-exposure was more photodamaging. Together, these data provide additional morphological evidence that AAF increased fibroblast photoresistance against both UVA and UVB.

The primary mechanism through which UVA induces cellular toxicity is through the generation of ROS [3, 9]; alternatively, UVB is cytotoxic primarily by directly damaging cellular DNA [2], though it too can generate ROS as a secondary mechanism of action [41].

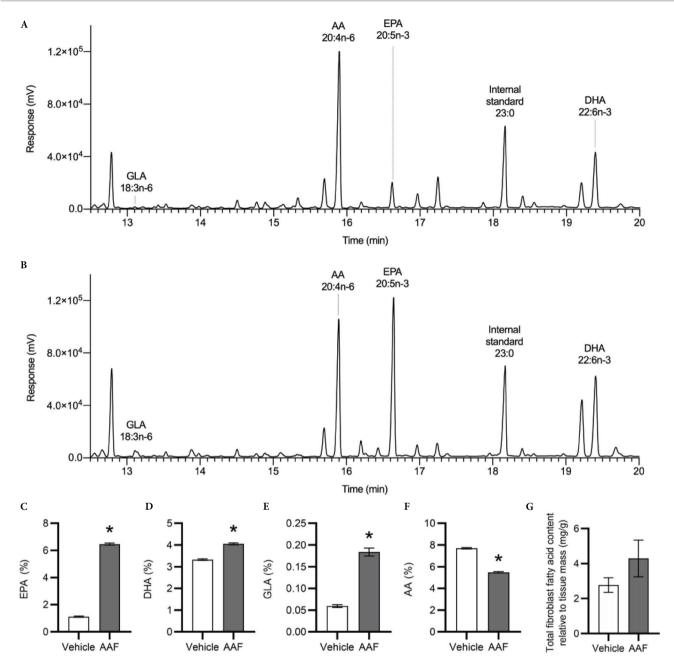


Figure 5. AAF alters lipid profile of the dermal fibroblasts. The fatty acid profile chromatograph examples depict how 14-day vehicle-treated dermal fibroblasts (A) have smaller response (mV) peaks for EPA, DHA, and GLA and a larger AA peak versus 14-day AAF-treated cells (B). Relative to total fatty acid content the mass percent of (C) EPA, (D) DHA, (E) GLA, and (F) AA, and (G) total fatty acid content relative to fibroblast mass are compared in 14-day vehicle- versus 14-day AAF-treated human primary dermal fibroblasts. \*Signifies value is significantly different from the vehicle control as determined by two-tailed t-tests (P < 0.05).

Based on the known antioxidant activity of AAF's carotenoid components, lutein and zeaxanthin [15, 16, 42], we hypothesized that one of its photoprotective mechanisms of action was through the induction of increased cellular antioxidant activity. However, due to incompatibilities between UV light and the ROS-detecting probe CM-DCFH<sub>2</sub>-DA often used in cellular antioxidant studies [43], we instead indirectly tested this hypothesis by determining if AAF-treatment induced a cytoprotective

effect against H<sub>2</sub>O<sub>2</sub>, a ROS induced by UV light in skin cells [41]. These experiments showed that AAF significantly reduced H2O2-induced cytotoxicity and intracellular ROS production with as little as 24 hours of treatment, supporting a rapid onset and cell-mediated antioxidant mechanism of action.

To explore the possibility that AAF was altering the activity of antioxidant pathways within the cells, we evaluated the effect of AAF on the expression of six



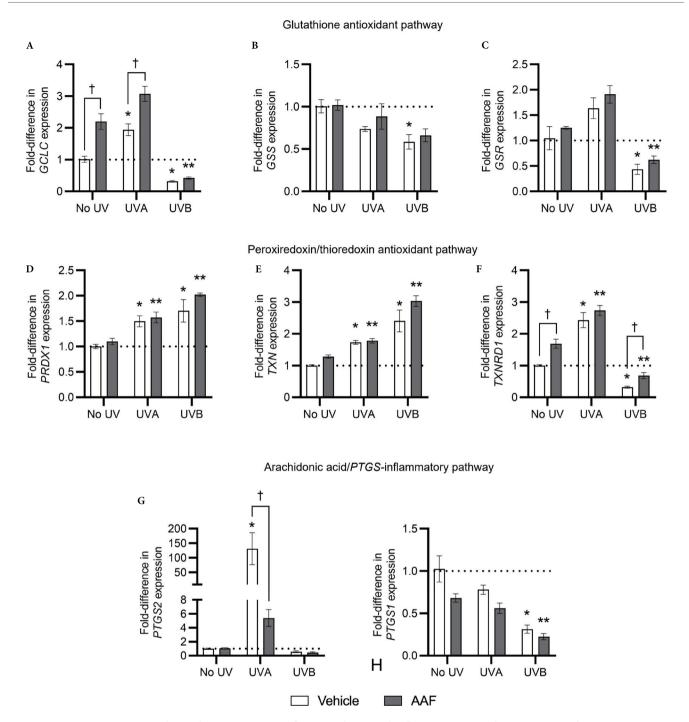


Figure 6. AAF treatment alters the expression of antioxidant and inflammatory pathway genes. The expression of genes involved in the glutathione antioxidant pathway, (A) GCLC, (B) GSS, and (C) GSR, the PRDX/TXN antioxidant pathway, (D) PRDX1, (E) TXN, and (F) TXNRD1, and the AA/PTGS-inflammatory pathway, (G) PTGS2 and (H) PTGS1, were quantified using qPCR on cDNA reverse transcribed from the isolated RNA of human primary dermal fibroblasts that had been treated with vehicle or AAF for 14 days, exposed to UVA (100 J/cm²) or UVB (1.5 J/cm²), and compared to baseline (No UV). \*Signifies value is significantly different from the vehicle-treated/baseline control, \*\*signifies value is significantly different from the AAF-treated/baseline control, and †signifies the two values indicated are significantly different from each other, as measured using two-way ANOVAs followed by Tukey's multiple comparisons test (P < 0.05) [34]. Statistical analyses were completed on the  $\Delta\Delta C_r$  values for each gene calculated via the  $2\Delta\Delta C_t$  method [35].

genes involved in the glutathione or PRDX/TXN antioxidant pathways. The antioxidant glutathione is biosynthesized in two steps that require the enzymes GCLC

[44–46] and GSS [47], with the activity of GCLC being the rate-limiting step [46]. A third enzyme, GSR, then reduces oxidized glutathione molecules to reactivate

their antioxidant capacity [48-50]. Our gene expression analysis suggests that AAF's UVA-protective effects may involve increased cellular antioxidant capacity via increased activity of the glutathione pathway, based on how it increased the expression of the rate-limiting GCLC. In the PRDX/TXN antioxidant pathway, PRDXs reduce H<sub>2</sub>O<sub>2</sub> to water, TXN reduces the PRDXs, TXNR reduces the TXN, and lastly nicotinamide adenine dinucleotide phosphate (NADPH) reduces the TXNR, thus reactivating the pathway [31, 51-53]. Our data suggest that this second photoprotective antioxidant pathway was upregulated by AAF treatment and, since AAF-treatment increased TXNRD1 expression in UVB-exposed fibroblasts, that this upregulation may be partly responsible for AAF's UVB-photoprotective properties. While our data provide initial support for an antioxidant photoprotective mechanism of AAF there are two key limitations: we did not directly measure intracellular ROS accumulation and its reversal by AAF in UV-treated cells, and we did not determine if the AAF-mediated changes in GCLC and TXNRD1 expression result in equivalent changes to the corresponding enzymes or their activity. Therefore, future experiments will be needed to confirm AAF's effects on these cellular antioxidant pathways in the setting of UVA and UVB exposures.

To further understand how AAF-treatment affected the fibroblasts, we performed a fatty acid analysis [54] which confirmed that EPA, DHA, and GLA, the LCPUFAs from AAF's fish and borage oil components, were all successfully integrating into the fibroblasts. A limitation of these trials is that we did not have the analytical capabilities to quantify the non-lipid components included in AAF: lutein, zeaxanthin, or vitamin D3. A key finding was that AAF diminished the mass percent of AA within the fibroblasts. Pro-inflammatory molecules called prostaglandins are biosynthesized from AA through PTGS enzymes [55], and since a major driver of UV-damage is inflammation [56], we hypothesized that AAF-treatment could dampen the inflammatory response through the AA/PTGS-inflammatory pathway as a second mechanism of photoprotection.

PTGS2 expression is induced during acute inflammatory events such as UV-exposure [57, 58], which catalyzes the first step in the conversion of AA to prostaglandins with prostaglandin E2 (PGE2) being the primary prostaglandin synthesized in the skin [59, 60], and is both overexpressed in and critical for the development of skin cancers [59, 61]. Consistent with the literature [59] we observed a robust increase in PTGS2 expression but no effect on the constitutively-expressed PTGS1 following exposure of fibroblasts to UVA. This

UVA-mediated increase in PTGS2 was largely abrogated by AAF (a 24.3-fold reduction). The reduction in cellular AA and PTGS2 expression suggests that AAF's UVA photoprotective mechanisms of action may also include a dampened acute inflammatory response via this pathway, though a limitation of our study is that we did not quantify PGE2 levels to confirm this. However, our data certainly warrants additional studies to confirm or rule out an anti-inflammatory photoprotective mechanism of AAF both in vitro and in vivo.

# Conclusions

This study shows that primary human dermal fibroblasts treated for 2 weeks with AAF are protected against the cytotoxic effects of UVA/B and ROS and suggest that AAF's photoprotective effects may include increased antioxidant capacity and a dampened inflammatory response. While more research is needed to confirm these putative mechanisms of action through enzyme activity-quantifying and in vivo assays, this study introduces valuable methodology for completing cellular photoprotection assays and is an important first preclinical step in determining how AAF induces its photoprotective effects.

# **List of Abbreviations**

AA: arachidonic acid

AAF: anti-aging formula

BHT: butylated hydroxytoluene

BSA: bovine serum albumin

BSO: borage seed oil

CM-DCFH<sub>3</sub>-DA: 5-(and 6-)chloromethyl-2'7'-dichlorodi-

hydrofluorescein diacetate DHA: docosahexaenoic acid

DMSO: dimethyl sulfoxide

EPA: eicosapentaenoic acid

FBS: fetal bovine serum

FO: fish oil

GC: gas chromatograph

GCFID: gas chromatography flame ionization detection

GCLC: glutamine-cysteine ligase catalytic subunit

GLA: gamma-linolenic acid

GSH: glutathione

GSR: glutathione reductase

GSS: glutathione synthetase

GSSG: glutathione disulfide

H<sub>2</sub>O<sub>2</sub>: hydrogen peroxide

IC<sub>50</sub>: half-maximal inhibitory concentration

LCPUFA: long chain polyunsaturated fatty acid

LSGS: low serum growth supplement

MED: minimal erythema dose



MFE: marigold flower extract

MTC: maximum tolerated concentration

MTC<sub>16</sub>: half of the maximum tolerated concentration

MTT: thiazolyl blue methyltetrazolium bromide

NADPH: nicotinamide adenine dinucleotide phosphate

PBS: phosphate buffered saline

PE: paprika extract PFA: paraformaldehyde PGE2: prostaglandin E2 PI: propidium iodide PRDX: peroxiredoxin PRDX1: peroxiredoxin 1

PTGS1/2: prostaglandin-endoperoxide synthase 1/2

qPCR: quantitative polymerase chain reaction

ROS: reactive oxygen species

THF: tetrahydrofuran TXN: thioredoxin

TXNRD1: thioredoxin reductase 1

UV: ultraviolet Vit D3: vitamin D3

## **Conflicts of Interest**

SRH was a recipient of a Mitacs Elevate Postdoctoral Fellowship award that was active from December 4, 2017 to December 3, 2019, and which was administered by KBG. The Award covered the salary for SRH and non-salary related research expenses for the study. Bend Beauty, Inc. contributed 60% of the total Mitacs award value. SRH then received direct salary support from Bend Beauty, Inc. from December 2019 to January 2020. AJR is an employee of and MSO the founder and president of Bend Beauty, Inc.

# **Ethics Approval and/or Participant** Consent

Our study did not require any ethics approval or participant consent because it was entirely completed using cell culture and in vitro experiments. No animal or clinical trials were completed for this study.

# **Authors' Contributions**

SRH: lead the design and planning of the study, collection, analysis, and interpretation of the data, drafted the manuscript, and gave final approval of the version to be published.

https://orcid.org/0000-0001-5314-4970

AJR: contributed to study design and planning, revised the manuscript critically, and gave final approval of the version to be published.

JE: collected and analyzed data, revised the manuscript critically, and gave final approval of the version to be published.

BTM: contributed to study design, revised the manuscript critically, and gave final approval of the version to be published.

MSO: contributed to study design and planning, revised the manuscript critically, and gave final approval of the version to be published.

KBG: contributed to study design and planning, analysis and interpretation of the data, revised the manuscript critically, and gave final approval of the version to be published.

https://orcid.org/0000-0003-0040-5879

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