Sulforaphane blocks hypoxia-mediated resistance to TRAIL-induced tumor cell death

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Abstract. Hypoxia occurs frequently in various solid tumors and elicits a cellular response designed to improve cell survival through adaptive processes, thereby accelerating cancer progression and the development of chemotherapy resistance. Tumor necrosis factor-related apoptosis-inducing ligand (TRAIL), a member of the TNF superfamily, leads to tumor cell death via both intrinsic and extrinsic apoptotic signaling pathways. Hypoxia inhibits TRAIL-mediated apoptosis and attenuates the therapeutic activity of TRAIL in cancer management. Hypoxia-inducible factor-1α (HIF-1α) plays a central role in tumor hypoxia by up-regulating gene expression related to angiogenesis, cancer invasion and anti-apoptosis. Sulforaphane (SFN), a phenethyl isothiocyanate, elicits HIF-1α inactivation under hypoxia. This study investigated whether hypoxic inhibition of TRAIL-mediated tumor cell death is increased by SFN-mediated HIF-1α instability. SFN induced cell death in various tumor cells, including SK-N-SH, SNU-638, HeLa and A549 cells, and showed cell cytotoxicity in hypoxia-exposed tumor cells. Western blot analysis showed that SFN treatment increased p53 and activated caspase-3 proteins, and decreased HIF-1α activation under hypoxia. Under low-oxygen conditions, TRAIL-treated cells displayed inhibited apoptosis, while SFN-pre-treated cells exhibited stronger sensitization to TRAIL under the hypoxic conditions. SFN treatment enhanced TRAIL-induced activation of proteins, including caspase-3 and p53. SFN dose-dependently decreased HIF-1α protein levels in cancer cells, which was mediated by decreased protein stability. This study demonstrated that SFN recovered hypoxia-mediated resistance to TRAIL via instability of HIF-1α, and also suggests that combination therapy with SFN and TRAIL may provide a novel strategy for treating hypoxic solid tumors.

Introduction

The programmed mode of cell death known as apoptosis can be triggered in tumor cells by anti-cancer agents (1,2). Tumor necrosis factor (TNF)-related apoptosis-inducing ligand (TRAIL), a member of the TNF family, is one of the stimuli that induces cancer cell death. TRAIL is of interest in oncology, since it displays specific anti-cancer activity against a wide range of cancer cells without significant side effects (3). In particular, some reports have shown that TRAIL induces apoptosis via a p53-mediated increase of DR5 (4,5). Additionally, one early study found that a number of tumor cell lines expressing wild-type p53 were significantly more sensitive to TRAIL as compared to mutations in p53 genes (6). Thus, TRAIL-mediated apoptosis has been related to p53 tumor suppressor functions during cancer therapy.

Hypoxia is a common stressor of tumors. In particular, rapidly-growing solid tumor cells are often hypoxic at their center. Hypoxia exhibits anti-apoptotic potential via dysregulation of a variety of apoptosis pathways, and decreases the effectiveness of anticancer drugs, including TRAIL, in solid tumor cells (7,8). Hypoxia-inducible factor-1 (HIF-1), a transcriptional factor composed of α- and β-subunits, is a key regulator of metabolic adaptation to hypoxia (9). HIF-1α expression is negatively regulated by the von Hippel-Lindau (pVHL) tumor suppressor gene. HIF-1α is hydroxylated at prolyl residues under normoxic conditions, thereby targeting the ubiquitin system (10). By contrast, under hypoxic conditions HIF-1α acts as the primary transcription factor functioning to activate multiple target genes (10). Targets of HIF-1α include cytokines and growth factors, angiogenesis-promoting genes, cell cycle progression, glucose uptake and metabolism, and cell survival. Thus, HIF-1α has been implicated as an oncogene that is overexpressed in human cancer cells (11). Appropriately, blockade of HIF-1α as a therapeutic target alone or in combination with chemotherapeutic reagents has been explored. HIF-1α has also been implicated as a co-regulator of transcription with p53 to promote apoptosis (12). Indeed, the apoptotic function of p53 can be regulated by the status of HIF-1α in cells, and blocking HIF-1α expression drives p53-mediated tumor cell death in hypoxia (13,14).

Sulforaphane (SFN), an isothiocyanate derived from the hydrolysis of glucoraphanin in broccoli, induces phase II
detoxification enzymes and inhibits chemically-induced mammary tumors in rodents (15). Recently, SFN was demonstrated to induce cell cycle arrest and apoptosis in human cancer cells (16). In addition, the compound is safe and essentially non-toxic, making it an excellent candidate as a chemoprevention agent. Recently, it was reported that SFN inhibits the proliferation of human cancer cells by inducing HIF-1α instability (17). However, the effect of the combination of SFN and TRAIL under hypoxic conditions has not been reported. The present study investigated whether SFN diminishes the hypoxic inhibition of TRAIL-induced apoptosis via the down-regulation of HIF-1α, and also indicates that SFN may act as a sensitizer for hypoxia-induced TRAIL-resistant cancer cells.

Materials and methods

Cell culture and hypoxic treatment. Human neuroblastoma cells (SK-N-SH), human uterine cervical cancer cells (HeLa), human alveolar basal epithelial cell (A549) and a human gastric cancer cell line (SNU 638) were obtained from the American Type Culture Collection (ATCC). Cells were cultured in Minimum Essential Medium (MEM; Hyclone) containing 10% fetal bovine serum (Gibco) and gentamycin (0.1 mg/ml) in a humidified incubator maintained at 37˚C and 5% CO₂. A hypoxia chamber was used to create a low-oxygen environment; a gas mixture of 1% O₂, 5% CO₂ and 94% N₂ flowed into the sealed chamber. Ambient air was evacuated by an outlet tube and O₂ flowed through the chamber for 2-3 min to maintain the desired O₂ tension.

Crystal violet assay. Cell viability was determined by crystal violet staining as previously described (18). Briefly, cells were stained with staining solution (0.5% crystal violet in 30% ethanol and 3% formaldehyde) for 10 min at room temperature (RT), washed four times with water and dried. Cells were then lysed with 1% sodium dodecyl sulfate (SDS) solution and cell absorbance was measured at 550 nm. Cell viability was calculated based on the relative dye intensity compared to the controls.

Terminal deoxynucleotidyl transferase dUTP nick end labeling (TUNEL) assay. TUNEL analysis was performed to measure the degree of cellular apoptosis using an in situ ApoBrdU DNA fragmentation assay kit (BioVision, CA, USA) following the manufacturer’s instructions.

DNA fragmentation assay. Cell pellets were suspended in 0.5 ml of 50 mM Tris-HCl buffer (pH 8.0) containing 10% (v/v) NP-40, 20 mM EDTA and 0.5% (w/v) SDS, and digested with 500 µg/ml proteinase K for 4 h at 65˚C. After digestion, the DNA was sequentially extracted with phenol/chloroform (1:1, v/v) and precipitated with ethanol at -20˚C overnight. Purified DNA was electrophoresed on a 1.5% agarose gel and visualized by ethidium bromide staining and ultraviolet transillumination.

Immunofluorescence. Cell lines cultured on glass coverslips were treated with SFN (Sigma) and/or TRAIL. Cells were washed with phosphate buffered saline (PBS) and fixed with cold acetone for 90 sec at RT. Cells were then washed with PBS again, blocked with 5% fetal bovine serum (FBS) in tris buffered saline with Tween (TBST) and incubated with anti-caspase-3 (2 µg/ml) and anti-HIF-1α (2 µg/ml) monoclonal antibodies for 48 h at RT. Unbound antibody was removed by an additional PBS wash and cells were incubated with labeled anti-rabbit Alexa Fluor 546 (for anti-caspase-3 and anti-HIF-1α) IgG antibodies (4 µg/ml) for 2 h at RT. Finally, cells were mounted with DakoCytonation fluorescent medium and visualized using fluorescence microscopy.

RT-PCR. Total RNA was isolated using RNAiso Plus (Takara). cDNA synthesis was performed using the Takara Prime Script™ first strand cDNA synthesis kit (200 U/µl) according to the manufacturer’s instructions. HIF-1α specific primer sequences (forward, 5’-AGAAACCACCTATGCCTGC-3’; reverse, 5’-GTGCGCTGAATAATACCCTC-3’) and β-actin-specific primer sequences (forward, 5’-GCAAGCGAGGTATGACGG-3’; reverse, 5’-CAAATAGCCATGGACCTTC-3’) were used for RT-PCR. Each 50 µl of amplification reaction contained 1.25 units of Taq polymerase, 2.5 mM dNTP mixture, 20 pM primers and 2 µl cDNA for each gene. The PCR parameters were 30 cycles as follows: denaturation at 94˚C for 30 sec, annealing at 55-60˚C for 30 sec, and extension at 72˚C for 1 min. The PCR products were separated on 1.2% agarose electrophoresis gels, stained with 0.5 µg/ml ethidium bromide and recorded for analysis.

Western blotting. Cell lines were lysed in a lysis buffer (25 mM HEPES; pH 7.4, 100 mM NaCl, 1 mM EDTA, 5 mM MgCl₂, 0.1 mM DTT and protease inhibitor mixture). Proteins were electrophoretically resolved on an 8-15% SDS gel, and immunoblotting was performed as previously described (19). Equal amounts of protein lysates were resolved on an 8-15% SDS-polyacrylamide gel and electrophoretically transferred to a nitrocellulose membrane. Immunoreactivity was detected through sequential incubation with horseradish peroxidase-conjugated secondary antibodies and ECL reagents. The antibodies used for immunoblotting were p53, HIF-1α and β-actin (Santa Cruz, CA, USA).

Statistical evaluation. All data are expressed as the means ± standard deviations (SD) and were compared using the Student’s t-test and the ANOVA Duncan test with the SAS statistical package. The results were considered significant for values of *P<0.05 or **P<0.01.

Results

SFN induces cancer cell apoptosis under normoxia or hypoxia. Hypoxia inhibits TRAIL-mediated apoptosis (8). On the other hand, SFN is an anti-cancer therapeutic agent that decreases HIF-1α activation under low-oxygen conditions (17). Appropriately, the present study examined the influence of hypoxia on SFN- or TRAIL-induced cell death in SK-N-SH, SNU-638, HeLa and A549 cancer cells. Crystal violet staining showed that all cell types were sensitive to SFN with or without hypoxia (40-60% reduction in viability of all cell populations) (Fig. 1A and C). On the other hand, hypoxia-exposed cells were resistant to TRAIL treatment.
To confirm whether SFN induced cell death, the morphology of SK-N-SH cells exposed to 20 µM SFN for 24 h and/or hypoxia was examined. Morphologic changes were observed in SFN-treated cells with or without hypoxia (Fig. 1B). Furthermore, SFN induced DNA fragmentation in SK-N-Sh cells under normoxia or hypoxia (Fig. 1D).

**SFN induces apoptotic signaling under hypoxic conditions through HIF-1α inactivation.** SFN decreases HIF-1α stability under hypoxia (17). Thus, immunocytochemistry was performed using an antibody against HIF-1α. Cells were treated with 20 µM SFN with or without hypoxia. The results showed that HIF-1α was detectable upon exposure to hypoxia and was decreased by SFN treatment (Fig. 2A). Since HIF-1α is known to negatively regulate p53 (14), the possible involvement in SFN-induced apoptosis was ascertained by immunoblot analysis using antibodies against HIF-1α and p53 (Fig. 2B). SFN-treated cells displayed increased expression of p53 protein under normoxia or hypoxia, while HIF-1α protein levels were decreased under hypoxia. These results are consistent with an effect of SFN on the expression of p53 protein levels via decreased HIF-1α protein expression under hypoxia. Analysis of mRNA expression showed that HIF-1α gene transcription was not affected by SFN under normoxia or hypoxia (Fig. 2C), indicating that SFN affected HIF-1α stabilization under hypoxia. Additionally, since SFN is a cancer therapeutic agent, immunocytochemistry analysis was performed using an antibody against the cleaved fragment of caspase-3, which is a key substrate for apoptosis. Cleaved caspase-3 was detectable in SFN-treated cells (Fig. 2D), indicating that SFN induced apoptosis through decreased HIF-1α stabilization and increased p53 expression and the activation of caspase-3 cleavage.

**SFN recovers hypoxic inhibition of TRAIL-mediated apoptosis by HIF-1α instability.** To determine whether HIF-1α instability is a mechanism of SFN-induced apoptosis under hypoxic conditions, an experiment was conducted in solid tumor cells to address the hypothesis that SFN-induced suppression of HIF-1α stabilization recovers TRAIL-induced apoptosis under hypoxia. This effect of SFN was observed in all cell lines (Fig. 3A). A reduction in cell viability of 40-50% was evident in SFN pre-treated cells subsequently exposed to TRAIL with or without hypoxia. The data was further corroborated by morphologic examination of hypoxic cells, which revealed a significant increase in the cell death of SFN and TRAIL combination-treated cells compared to cells treated only with TRAIL (Fig. 3B). Treatment with SFN and TRAIL increased p53 and decreased HIF-1α protein expression under hypoxia (Fig. 4A). Immunofluorescent analysis using an antibody against cleaved caspase-3 detected the cleavage product in SK-N-SH cells exposed to SFN and TRAIL under hypoxic conditions (Fig. 4B). The results of a TUNEL assay also showed elevated sensitivity to TRAIL protein in hypoxic tumor cells (Fig. 4C). The results indicate that HIF-1α activity suppressed by SFN sensitized hypoxic tumor cells to TRAIL-induced apoptosis.

**Discussion**

The main focus of this study was the investigation of the therapeutic potential of the HIF-1α suppressor compound SFN...
in therapeutic intervention strategies using TRAIL protein against hypoxic solid tumors.

In response to hypoxia, cells alter their metabolism and regulation of gene expression to adapt to hypoxia (9). Previous studies have shown that hypoxic tumor cells block many different therapies through the increased expression of HIF-1α, Bcl-2 and IAP family members, while decreasing the expression of the p53 and caspase family (7,20). Consistent with these results, in the present study the expression of p53 protein was decreased and the expression of HIF-1α was
Figure 4. SFN recovers TRAIL-mediated apoptosis under low-oxygen conditions via HIF-1α instability. (A) Western blot analysis of HIF-1α and p53 in SK-N-SH cells. Cells were pre-incubated under hypoxic conditions for 12 h, and then exposed to 10 μM SFN for 12 h in the absence of TRAIL or presence of 400 ng/ml TRAIL for 6 h. β-actin served as the loading control. (B) Immunocytochemistry of cleaved caspase-3 in SK-N-SH cells. Cells were pre-incubated under hypoxic conditions for 12 h, and then exposed to 10 μM SFN for 12 h in the absence of TRAIL or presence of 400 ng/ml TRAIL for 6 h. Scale bar, 100 μm. (C) Representative immunofluorescence images of TUNEL-positive (white) SK-N-SH cells pre-incubated with hypoxia and then with 10 μM SFN for 12 h in the absence of TRAIL or presence of 400 ng/ml TRAIL. The cells were counterstained with propidium iodide (grey) to show all cell nuclei. Scale bar, 20 μm. *P<0.01, **P<0.05, significant differences between the control and each treatment group.

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References


