Mitochondrial DNA depletion causes decreased ROS production and resistance to apoptosis

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Abstract. Mitochondrial DNA (mtDNA) depletion occurs frequently in many diseases including cancer. The present study was designed in order to examine the hypothesis that mtDNA-depleted cells are resistant to apoptosis and to explore the possible mechanisms responsible for this effect. Parental human osteosarcoma 143B cells and mtDNA-deficient (Rho˚ or q°) 206 cells (derived from 143B cells) were exposed to different doses of solar-simulated ultraviolet (UV) radiation. The effects of solar irradiation on cell morphology were observed under both light and fluorescence microscopes. Furthermore, apoptosis, mitochondrial membrane potential (MMP) disruption and reactive oxygen species (ROS) production were detected and measured by flow cytometry. In both cell lines, apoptosis and ROS production were clearly increased, whereas MMP was slightly decreased. However, apoptosis and ROS production were reduced in the Rho˚206 cells compared with the 143B cells. Also performed western blot analysis and demonstrated the increased release of cytosolic Cyt c from mitochondria in the 143B cells compared with that in the Rho˚206 cells. Thus, we concluded that Rho˚206 cells exhibit more resistance to solar-simulated UV radiation-induced apoptosis at certain doses than 143B cells and this is possibly due to decreased ROS production.

Introduction

Mitochondrial DNA (mtDNA) is comprised of a circular molecule of approximately 16.6 kb, encoding RNAs and polypeptides of the mitochondrial respiratory chain. Intact mtDNA is necessary for the production of these polypeptides, which consist of the key catalytic subunits of the mitochondrial respiratory chain complexes and are essential for oxidative ATP production. The mtDNA is frequently exposed to oxidative stress due to the process of oxidative phosphorylation. During oxidative phosphorylation, molecular oxygen is transformed into highly reactive oxygen species (ROS) in the electron transport chain, resulting in the damage of functional macromolecules including DNA (1).

The occurrence of mtDNA variation during evolution, particularly with regard to the normal variation in its amount, is closely associated with cell physiology and human health. Over the past 20 years, researchers have found that a number of disorders in humans are associated with severe mtDNA depletion and that long-term exposure to low concentrations of ethidium bromide (EB) establishes mtDNA-deficient (Rho˚ or q°) cells (2). Thus, q° cells have been used to examine the roles of mtDNA and particularly the importance of mtDNA in apoptosis (3). A previous study showed that respiration-deficient cells resisted tumor necrosis factor (TNF)/serum deprivation-induced apoptosis, whereas apoptosis was initiated in parental cells and rescued cells with normal mtDNA (4). These findings suggest that mtDNA depletion is involved in a mechanism responsible for resistance to apoptosis. Furthermore, studies by Amuthan et al and Biswas et al demonstrated that mtDNA depletion contributed to tumor progression and metastasis (5,6). Thus, it is likely that mtDNA depletion prevents apoptosis and generates cancer-related proteins.

Organisms may be exposed to numerous noxious agents under various conditions. These agents not only passively disintegrate cells, but also induce productive responses. In particular, it has been shown in mammalian cells that several genes are activated by ultraviolet (UV) irradiation (7).

Taking all of the above into consideration, this study was designed to examine the hypothesis that mtDNA-depleted mammalian cells resist UV-induced apoptosis and to explore the possible mechanism responsible for this effect.

Materials and methods

Cell culture, reagents and antibodies. The human parental osteosarcoma cell line 143B and Rho˚206 cells
Table I. UVA + UVB intensity, exposure time and dosage.

| Experimental grouping         | Intensity (UVA + UVB mW/cm²) | Exposure time (min) | Dosage (mJ/cm²+J/cm²) |
|-------------------------------|-----------------------------|--------------------|----------------------|
| Sham-irradiation group (SIG)  | 0+0                         | 0                  | 0+0                  |
| Low-dose groups (LDG)         |                             |                    |                      |
| IG1                           | 2.95+56.5                   | 1                  | 177+3.39             |
| IG2                           | 2.95+56.5                   | 2                  | 354+6.78             |
| High-dose groups (HDG)        |                             |                    |                      |
| IG3                           | 2.95+56.5                   | 3                  | 531+10.17            |
| IG4                           | 2.95+56.5                   | 4                  | 708+13.56            |

(mtDNA-depleted) were a gift from Professor Minxin Guan (Zhejiang University, Zhejiang, China). The cells were cultured in Dulbecco's modified Eagle's medium (Invitrogen, Carlsbad, CA, USA) supplemented with 10% newborn calf serum (Gibco, Carlsbad, CA, USA), 100 U/ml penicillin, 100 mg/ml streptomycin, 100 µg/ml bromodeoxyuridine and 50 µg/ml uridine (Sigma-Aldrich, St. Louis, MO, USA). Rhodamine 123 (Rh123) and dichlorodihydrofluorescein diacetate (DCFH-DA) were purchased from Sigma Chemical Co. (St. Louis, MO, USA). 3-(4,5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) was purchased from Promega (Madison, WI, USA). We purchased cytochrome c (Cyt c) antibodies (12963S) from Cell Signaling Technology (CST; Beverly, MA, USA). Antibodies against β-actin (AA128) and the secondary horseradish peroxidase (HRP)-labeled antibodies (A0216), were purchased from Beyotime Biotechnology (Jiangsu, China).

Irradiation procedure. The irradiation procedure was performed using a 1000 Watt Solar Oriel UV Simulator (Oriel, Stratford, CT, USA) with a UVX digital radiometer (Ultra-Violet Products, Upland, CA, USA) which was equipped with a UVX-310 sensor to measure UV radiation intensity. The intensities of UVA and UVB were 2.95 and 56.6 mW/cm², respectively, with doses ranging from 177 mJ/cm² and 3.39 J/cm² to 708 mJ/cm² and 13.56 J/cm², respectively (Table I). The cells were washed twice with phosphate-buffered saline (PBS) prior to UV irradiation and then covered with a thin film of PBS during UV irradiation. Following irradiation, PBS was removed and immediately replaced with the maintenance medium. The sham-irradiated cells (control group) were similarly treated; however, they were exposed to normal room lighting. Finally, all the cells were incubated at 37°C in an incubator with 5% CO₂ (Thermo Fisher Scientific, Waltham, MA, USA).

Cell viability assay. As previously described, the MTT method (8) was performed to measure cell viability. Prior to adding MTT tetrazolium salt to each well at a working concentration of 5 mg/ml, the cells were seeded in 96-well plates at a density of 5×10⁴ cells/cm² overnight and treated with UV as indicated in Table 1 with a 4 h incubation time in a CO₂ incubator. Subsequently, the medium in each well was replaced with 150 µl DMSO (Sigma-Aldrich). The absorbance of the dissolved formazan crystals in each well was measured using a plate reader (Dynatech MR5000; Dynex Technologies, Chantilly, VA, USA) at a test wavelength of 570 nm. Cell viability was calculated using the following equation: (absorbance of the experiment samples/absorbance of the control) x100%.

Measurement of mitochondrial membrane potential (MMP). The cationic lipophilic green fluorochrome Rh123 is an effective reagent used to determine MMP (9). Prior to incubating with 10 µM Rh123 at 37°C for 30 min, the cells were harvested and washed three times with PBS. Following incubation, the cells were washed twice with PBS and fluorescence was determined using a flow cytometer (Beckman Coulter, Inc., Brea, CA, USA) with an excitation wavelength of 488 nm through the FL-1 filter.

 Determination of intracellular ROS levels. DCFH-DA is a reliable reagent which was used to determine intracellular ROS levels as previously described (10). The cells were incubated with DCFH-DA (10 µM) at 37°C for 30 min. Following incubation, the cells were washed with PBS twice and the fluorescence of dichlorofluorescein (DCF) generated by ROS was analyzed immediately using a flow cytometer through the FL-1 filter with an excitation wavelength of 488 nm.

Annexin V-PE/7-AAD staining. Apoptosis was determined by flow cytometric analysis (BD Biosciences, Franklin Lakes, NJ, USA) with Annexin V-PE and 7-AAD double staining. After the cells were harvested and washed twice with ice-cold PBS, the cells were stained and fluorescence was measured at an excitation wavelength of 480 nm through the FL-1 filter (530 nm) and FL-2 filter (585 nm).

Assessment of cell morphology and PCR identification. Cell morphology was assessed using a fluorescence microscope and an electron microscope (Jeol, Peabody, MA, USA). For microscope observations, the cells were firstly observed under bright light after being cultured overnight in 24-well dishes and exposed to UV radiation as described in Table I, and subsequently observed under purple light after fixing and staining with DAPI (Beyotime Biotechnology) for 15 min.

For electron microscopy, the cells were fixed with 2% paraformaldehyde/2% glutaraldehyde in 0.1 M PBS,
followed by 1% OsO4 (Sigma-Aldrich). Finally, the cells were
stained with uranyl acetate (Sigma-Aldrich) and lead citrate
(Sigma-Aldrich) for observation after dehydration, as previ-
ously described (11).
A volume of 20 µl containing 10 µl PrimeStar Max Premix 2X (Takara, Dalian, China) and 10 pmol of primer
was added to amplify the COX II gene. The human mito-
chondrial COX II primers used in our study have been reported
previously (12). The following sequences of COX II primers
were used: forward, 5'-ATC AAA TCA ATT GGC CAC CAA
TGG TA-3' and reverse, 5'-TTG ACC GTA GTA TAC CCC
CGG TC-3' (297 bp). PCR was performed according to the
manufacturer's instructions. The 200bp DNA Ladder (Dye
Plus; Takara Bio, Otsu, Japan) was used to verify the bands.

Western blot analysis. The cells, at a density of 1x10^7 cells/ml,
were exposed to UV radiation as described in Table I. Following
the lysis procedure, the lysates were centrifuged to obtain
the supernatants. The protein concentrations of the super-
натants were then determined using BCA Protein assay
reagent (Beyotime Biotechnology). Western blot analysis
was performed to analyze certain proteins in the supernatants
(50 µg) of each sample.

Statistical analysis. Data are expressed as the means ± SD and
analyzed using the Student's t-test (two-tailed). A P-value <0.05
was considered to indicate a statistically significant difference.

Results

Rho^206 cells exhibit greater resistance to apoptosis
compared with 143B cells following exposure to UV radia-
tion. Previous studies have shown that transient mtDNA
depletion enhances the invasive ability of several types of
cancer cells as well as resistance to apoptosis (5,13). Thus,
it is evident that mtDNA-depleted cells play roles in tumor
progression and metastasis. To the best of our knowledge,
there are no studies regarding the effect of mtDNA deple-
tion in human osteosarcoma 143B cells. To determine the
anti-apoptotic effects of mtDNA depletion in 143B cells, we
used Rho^206 cells, an mtDNA-depleted 143B cell type, as
shown in Fig. 1.

Firstly, we observed the decrease in cell numbers in the
cell lines following exposure to UV radiation as described in
Table I under a light microscope and a fluorescence micro-
scope. Fig. 2A and B show that the numbers of 143B cells
clearly decreased in comparison with the number of Rho^206
cells. Furthermore, MTT analysis showed the differences
in the cell viability of Rho^206 and 143B cells following
exposure to UV radiation. The cell viability of 143B cells
was lower than that of Rho^206 cells (Fig. 2C). Additionally,
Annexin V-PE and 7-AAD double staining by flow cytometry
shows that the 143B cells endured more severe apoptosis
compared with the Rho^206 cells following exposure to UV
radiation (Fig. 2D and E). This may indicate that Rho^206 cells
are more resistant to apoptosis.

UV-induced disruption of MMP is reduced in Rho^206 cells
compared with that in 143B cells. To detect changes in MMP
in the cells following UV irradiation in the present study, we
measured MMP by assessing the uptake of Rh123 using a flow
cytometer.

Following exposure to UV radiation as described in
Table I, mitochondrial activity was decreased in both cell
lines. As shown in Fig. 3, following exposure to different
intensities of radiation, the 143B cells displayed various
disruptions of MMP. Compared with the control, mitochon-
drial activity decreased to 76.35% after 708 mJ/cm^2 UVA and
13.56 J/cm^2 UVB treatment (IG4 group) in the 143B cells, and
to 85.1% in the Rho^206 cells. The impact of the combination
of 708 mJ/cm^2 UVA and 13.56 J/cm^2 UVB on mitochondrial
activity in the 143B cells (Fig. 3B and C) suggested that UV
causen the decline in mitochondrial activity. At the same
time, no significant disruption of MMP was detected after
708 mJ/cm^2 UVA and 13.56 J/cm^2 UVB treatment in both cell
lines (data not shown), indicating that UV may exert a minimal
effect on the MMP and mitochondrial dysfunction.
Furthermore, Fig. 3 shows clearly that the MMP in the
143B cells dropped more quickly than in the Rho^206 cells
following UV irradiation from which we concluded that
mtDNA depletion may protect cells against UV-induced
disruption of MMP.

UV irradiation induces less ROS production in Rho^206
cells compared with that in 143B cells. Disruptions in MMP
and cell apoptosis are associated with the production of
ROS (14,15). Thus, in the present study, we measured ROS
production in the 143B cells and the Rho^206 cells following
exposure to UV radiation as described in Table I. The
oxidation sensitive fluorescent dye DCFH-DA was applied
in order to evaluate ROS production. ROS production was
significantly increased in both cell lines following exposure
to increasing doses of UV radiation. In the 143B cells, the
generation of ROS was faster compared with that in the

![Figure 1. Verification of mitochondrial DNA (mtDNA)-depletion in Rho^206 cells. The mtDNA-encoded gene COX II was amplified by PCR. A 297 bp band containing the COX II sequence was detected in the DNA isolated from 143B cells whereas it was lost in the DNA isolated from Rho^206 cells.](image-url)
Figure 2. Ultraviolet (UV) irradiation induces various degrees of cell death in 143B and Rho⁺206 cells. (A) Effect of solar-simulated UV irradiation on the morphology of 143B cells observed under a fluorescence microscope (x40 magnification). (B) Morphological changes showed in Rho⁺206 cells following treatment with solar-simulated UV irradiation under a fluorescence microscope (x40). (C) MTT analysis of 143B and Rho⁺206 cells following solar-simulated UV irradiation. *P<0.05 vs. SIG control. Each bar represents the means ± SD from three experiments. Annexin V-PE and 7-AAD staining for apoptosis in (D) 143B and (E) Rho⁺206 cells following simulated solar irradiation. x-axis, Annexin V-PE; y-axis, DNA content by 7-AAD. The experiment was repeated three times and the image presented is typical of these three independent tests.
control group after UV exposure, (an approximate 3-5-fold increase), as shown in Fig. 4A and C. In the Rho’206 cells, there was an approximate 2-4-fold increase in the generation of ROS compared with the control group (Fig. 4B and C). These results show that ROS production increased more rapidly in 143B cells compared with that in Rho’206 cells following UV irradiation.

Rho’206 cells resist UV-induced apoptosis through decreasing Cyt c release. During apoptosis, Cyt c is released from mitochondria into the cytoplasm (16). To examine whether Cyt c was released during UVA combined with UVB-induced cell apoptosis, the expression of cytosolic Cyt c was measured using western blot analysis. Increased levels of Cyt c in the cytoplasm were detected following exposure to a combination of UVA and UVB (Fig. 5) compared with the controls. This increase was more evident in the parental 143B cells than in the mtDNA-depleted Rho’206 cells, indicating a potential resistance to Cyt c release in the Rho’206 cells. Furthermore, the morphological changes observed in the mitochondria of the Rho’206 cells (Fig. 6) may not be conducive to Cyt c release in these cells.

**Discussion**

Previous findings have demonstrated that mtDNA encodes pivotal catalytic subunits and its complete depletion in cells leads to several morphological changes. Fig. 6 shows that the mitochondrial density of mtDNA-depleted Rho’206 cells was reduced and that the mitochondria were larger and more elongated compared with those in the parental 143B cells. Hojo et al also observed similar changes in cyclosporin-treated cells (17). Furthermore, a major block also occurs in the normal electron flow in mtDNA-depleted cells.

Mitochondria are the principal source of intracellular ROS under adverse conditions (18,19). This, in combination with the fact that excision repair frequently occurs in mtDNA (20), suggests that mtDNA is liable to attack under conditions of oxidative and chemical stress. A number of studies regarding homoplasmic/heteroplasmic mtDNA depletion/mutations have
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been reported in human tumors which support this view (21-27). However, it remains unclear whether cancer progression results from (28) or from (29) the depletion/mutations. The results presented in this study suggest that mtDNA-depleted cells possess a survival advantage following environmental exposure to UV irradiation.

UV irradiation is a pivotal factor that increases levels of ROS (30-35), and simultaneously decreases antioxidant enzymes (36), causing oxidative stress to initiate cellular signal transduction. Redundant ROS are likely to lead to cell death by oxidizing and then damaging functional macromolecules such as DNA and protein. Herein, we found that UV irradiation increases the production of reactive oxygen species (ROS) in human tumors, which supports this view (21-27). However, it remains unclear whether cancer progression results from the depletion/mutations. The results presented in this study suggest that mtDNA-depleted cells possess a survival advantage following environmental exposure to UV irradiation.

Figure 4. Exposure to ultraviolet (UV) radiation increases the production of reactive oxygen species (ROS). (A) 143B and (B) Rho°206 cells were seeded at a density of 1x10^6 cells/ml, and then exposed to different doses of UV (Table I). After three washes, the cells were incubated at 37°C for 30 min with 10 μM DCFH-DA in PBS. The fluorescence was measured using flow cytometry (excitation wavelengths were both 488 nm, and emission wavelengths were both 525 nm). (C) A comparison between 143B and Rho°206 cell lines shows that there was no significant difference in ROS production between the SIG groups (P>0.05). However, there were significant differences between the cell lines at the same irradiation doses. *P<0.05 vs. SIG control. The test was repeated three times and the image presented is representative of these three independent tests.

Figure 5. Effects of ultraviolet (UV) irradiation on the levels of cytochrome c (Cyt c) in the two cell lines. (A) Rho°206 cells and (B) 143B were exposed to different doses of UV radiation (Table I). The cytosolic fractions of Cyt c were measured using western blot analysis. The experiment was repeated three times and the image presented is representative of these three independent tests.
exposure induced ROS production in human mammalian cells and mtDNA-depleted cells resisted the release of ROS thereby escaping greater death compared with the normal cells. Notably, UV irradiation in 143B cells and Rho°206 cells had minor effects on MMP, since MMP disruption is always followed by increasing ROS levels (37). This paradox warrants further investigation. We also observed Cyt c release, as previously described (37). Notably, the 143B cells released more Cyt c from the mitochondria into the cytoplasm than the Rho°206 cells following UV irradiation. Therefore, it is possible that DNA depletion in mitochondria induces resistance to UV-induced apoptosis by decreasing ROS production, which plays a role in the upstream apoptotic mechanism following Cyt c release and the activation of caspases.

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