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# **Nucleophosmin Redistribution Following Heat Shock: a Role in Heat-Induced Radiosensitization**

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## **Abstract**

Cellular survival from radiation induced DNA damage requires access to sites of damage for the assembly of repair complexes and the subsequent repair, particularly the repair of DNA double strand breaks (DSBs). Hyperthermia causes changes in protein-protein/DNA interactions in the nucleus that block access to sites of DNA damage. Studies presented here indicate that the nucleolar protein, nucleophosmin (NPM), redistributes from the nucleolus following hyperthermia, increases its association with DNA, and blocks access to DNA DSBs. Reduction of NPM significantly reduces heat-induced radiosensitization, but reduced NPM level does not alter radiation sensitivity *per se*. NPM knock down reduces heat-induced inhibition of DNA DSB repair. Also, these results suggest that NPM associates with nuclear matrix attachment region (MAR) DNA in heat shocked cells.

#### **Keywords**

Nucleophosmin; Hyperthermia; Radiosensitization

# **INTRODUCTION**

The potential for using hyperthermia as a radiosensitizing adjuvant for radiation therapy has long been recognized. Recent meta-analysis shows that hyperthermia significantly improves radiation therapy (1). Further, hyperthermia is a potent radiosensitizer, producing thermal enhancement ratios (TERs) from 3 to 5 (1,2). Nevertheless its clinical use remains limited, in part, because the mechanism of hyperthermia induced radiosensitization remains poorly understood. Reasonably, inhibition of DNA double strand break (DSB) repair should be involved in heat-induced radiosensitization. However, the literature indicates a more complex situation. Cells defective in the non-homologous end joining pathway or in homologous recombination show similar TERs but are more radiosensitive than isogenic wild type cells (2). One interpretation of these results is that neither of these pathways is a critical target for heat-induced radiosensitization. However, since the mutants studied are defective in the effector mechanisms of these pathways, an alternative interpretation is that hyperthermia affects an upstream step such as the sensing of DNA damage which would impact both DNA DSB repair pathways. One explanation  $(3-6)$  suggests that masking of DNA strand breaks by heat-induced protein-DNA complexes would sterically inhibit access of the sensing proteins to the damage site. However, little work has been done toward determining what protein(s) involved in DNA damage masking and the DNA regions masked in heat shocked cells. Here

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we present evidence that heat-induced masking of DNA damage is due (in part) to the heatinduced association of the nucleolar protein, nucleophosmin (NPM) with MAR DNA.

NPM is an abundant phosphoprotein, localized to the granular regions of the nucleolus (7). NPM trafficking contributes to its essential functions (8). Thus, changing the ability of NPM to mobilize to distinct cellular locales should alter NPM function (8). Additionally, loss of NPM function results in attenuation of cellular proliferation and increased apoptosis (9-16). The nucleolus acts as repository of stress responsive proteins (17-20). Specifically, the nucleolar protein, nucleolin (NCL or C23), redistributes from the nucleolus and interacts with replication protein A resulting in inhibition of DNA synthesis (20-22). The nucleolus is a heat sensitive organelle, undergoing detectable changes after 1 hr at 41°C and general changes after 15 min at 42°C, which leave cytoplasmic organelles largely unaffected (23-25). Further, HSP70 and HSP110, which modulate heat-induced radiosensitization localize in the nucleolus in cells during and immediately after heat shock (26-27). Therefore, we determined if NPM could play a key role in heat radiosensitization.

### **MATERIALS AND METHODS**

#### **Cell Lines and Cell Culture**

HeLaS3 cells (ATCC) were maintained in alpha-MEM media with 10% calf serum, 100 U/ml penicillin and 100 mg/ml streptomycin. Human breast cancer cells, HCC1806, were maintained in RPMI 1640 media with 10% Bovine Growth Serum and antibiotics. For survival experiments, control and treated cells were plated in replicate T-25 flasks. After 10 days (HeLaS3) or 19 days (HCC1806), flasks were stained with Crystal Violet and viable colonies  $(\geq 50 \text{ cells})$  counted.

#### **Hyperthermia and Irradiation**

For acute hyperthermia, flasks were immersed in a temperature controlled water bath. Irradiation was performed using a PANTAK pmc1000 X-ray machine with a  $0.1 \text{ Cu}^+ 2.5 \text{ AL}$ filter at a dose rate of 88.7 cGy/min.

#### **Transfection with siRNA**

HeLaS3 cell transfections were accomplished using a Nucleofector II electroporation apparatus (Amaxa) following the manufacturer's instructions. NPM specific siRNA sequences (28) and siCONTROL Non-Targeting siRNA #1 were obtained from Dharmacon. GFP-NPM (29) images were captured with a Spot Pursuit digital camera (Diagnostic Instruments) on an Olympus fluorescent microscope.

#### **Assay for DNA Double Strand Breaks**

The Neutral Comet Assay was performed using the Comet Assay Kit from Trevigen following manufacturer's instructions. Proteinase K (Sigma), when used, was added to the lysis solution (0.27 U/ml). Lysis occurred at 4°C for 30 min. Comets were analyzed by CometScore software (TriTek).

#### **Assay for DNA Supercoiling Ability**

The halo assay was done as described in (30).

#### **Chromatin Immunoprecipitation (CHiP) Assay**

The enzymatic shearing Chip-IT kit (Active Motif) was used following the manufacturer's instructions. NPM was immunoprecipitated with mouse anti-NPM antibodies (Zymed

Invitrogen). Co-immunoprecipitated DNA fragments were PCR amplified using primers to Topoisomerase I; MAR II, Exon 13, Exon 20; c-jun and the GAPDH promoters (31).

#### **Immunoblots, Antibodies and Other Reagents**

SDS-PAGE and immunoblotting were done following standard procedures using a BioRad Mini-Protean II gel and blotting system. Immunoblots were probed with mouse anti-NPM or HSP70/HSC70 antibodies and alkaline phosphatase conjugated Affinipure Goat anti-Mouse IgG+IgM secondary antibodies. The targeted bands were visualized colorimetrically by BCIP/ NBT.

# **RESULTS**

To confirm that heat increased NPM-DNA binding we crosslinked DNA-associated proteins to DNA with cisplatin, then isolated the DNA with hydroxyapatite, and resolved the DNA associated proteins by 2D gel electrophoresis. Hyperthermia increased the DNA association of NPM isoform, B23.1, more than 6-fold. Several NPM isoforms became associated with DNA in heated cells (Fig. 1A). These results are consistent with previous reports showing that hyperthermia increases NPM association with DNA in vivo (5) binds DNA in vitro (32).

To determine the effects of hyperthermia on the sub-nuclear localization of NPM we exposed HeLaS3 cells and HCC1806 cells to increasing heat shocks and visualized the localization of NPM using fluorescently labeled antibodies or transiently expressed GFP-tagged NPM fusion protein. NPM readily re-localizes from the nucleolus to the nucleoplasm after heat shocks of 15-45 min at 45°C in HeLaS3 and HCC1806 cells (Fig. 1). Thus, NPM redistributes into the nucleoplasm after hyperthermia and becomes associated with DNA.

Since NPM becomes associated with DNA, we determined if NPM could be involved in radiation sensitivity. We measured the TER after a 90-95% NPM knockdown, using siRNA. Cells were X-irradiated with or without prior hyperthermia (30 min at 45°C) and their colony forming ability measured (Fig. 2A). For HeLaS3 and HCC1806 cells, TER values were calculated at the control survival level after 2 and 4 Gy which were chosen because these doses are commonly used in clinical radiation therapy and averaged. Hyperthermia significantly increased the non-transfected cell's sensitivity to X-rays, TER = 2.22 and 2.08 for HeLaS3 and HCC1806, respectively (Fig. 2D). Transfection with the nonspecific siRNA did not significantly change the TER (2.22 and 2.16). However, transfection with the siRNA specific for NPM significantly lowered the TER to 1.74 and 1.39 (p<0.05; Students t-test) in HeLaS3 and HCC1806 cells, respectively. Also, NPM reduction made a statistically significant increase in survival at each dose level by the Student's t-test. Thus, heat induced radiosensitization is due, in part, to the presence of NPM. Without heat shock, NPM knockdown had no effect on survival after 2, 4 or 6 Gy (Fig. 2A).

Transient NPM knockdown and transfection with a non-specific siRNA did not significantly alter the plating efficiency in HeLaS3 cells  $(80 \pm 3\%)$  or in HCC1806 cells  $(30 \pm 3\%)$ . However, NPM knockdown sensitized HeLaS3 cells to heat shock alone. To understand this observation, we measured HSC70/HSP70 levels. NPM knockdown reduced the endogenous levels of HSC70/HSP70 (Fig. 2C), which confers resistance to heat-induced cell killing (33-34). This result also supports the hypothesis that NPM specifically contributes to heat induced radiosensitization; because its knockdown increased thermal sensitivity while reducing the TER without affecting radiation sensitivity *per se*. In contrast, nonspecific thermal sensitizers and protectors have been shown to alter both heat sensitivity and the TER in concert (35).

Since NPM knockdown reduced the TER from heat shock, we determined if heat induced a NPM dependent masking of DNA DSB and the inhibition of DNA DSB repair. Using the

neutral comet assay with PK (0.27 U/ml) the level of DNA damage immediately after 8 Gy, was the same between cells heated at 45<sup>o</sup>C for 30 min and unheated. However, a significant inhibition of subsequent DNA DSB repair could be observed (Fig. 3A, **left**). In control cells 75% of the DNA DSBs were repaired after 90 min, but in heated cells only 20-25% was repaired. After 150 min, more than 90% of the DNA DSBs were repaired in unheated cells, while  $\sim$  50% remained unrepaired in the heated cells. In the absence of PK there is a 60% reduction in the measured comet moment (Fig. 3A, **right**) for cells, heated at 45°C for 30 min, compared with unheated cells. Since the comet moment is the same in heated versus non-heated cells in the presence of PK, we conclude that 60% of the DNA DSBs, following 8 Gy, are masked (Fig. 3A, **right**), by a heat-induced, PK digestible component i.e. a protein or proteins. We define masking as the inability of an assay (or DNA repair pathway) to detect DNA damage in the presence of protein-DNA interactions even though the damage would be detectable in the absence of such protein-DNA interactions. Without PK, there appeared to be no repair of DNA DSB for 150 min (Fig 3A **left**), which may be due to a combination of unmasking the DSBs and their subsequent repair.

We determined the effects of NPM knockdown on the heat-induced masking of DNA damage and repair inhibition. The masking of DNA DSB is reduced by 40% in cells treated with the NPM siRNA (Fig. 3B **right**). To illustrate the effects of NPM knockdown on the masking of DNA damage, the comet moment values immediately after irradiation are plotted (Fig. 3C). Hyperthermia reduced the post 8 Gy comet moment ~3-fold. The comet moment was reduced  $\sim$ 2-fold after NPM knockdown (p < 0.05). Heated NPM knockdown cells, 90 min after 8 Gy, had 22% of the initial DNA DSB remaining for a 5-fold reduction in repair inhibition. Further the residual unrepaired DSB was reduced from 50% unrepaired to 12% unrepaired. Unrepaired DSBs are critical in hyperthermic radiosensitization (36-37). In contrast, in cells treated with a nonsense siRNA the heat-induced masking of DNA DSB and inhibition of their repair were identical to the untreated control (Fig. 3B **right**). Without heat shock, NPM knockdown had no effect on DNA DSB repair (Fig. 3B **left**) consistent with the observation that NPM knockdown had no effect on radiosensitivity in the absence of heat shock.

Previous studies of the masking effect using the halo assay, a DNA supercoiling assay that depends on DNA-nuclear matrix anchoring (38) found that it correlated with the TER in cell lines having various levels of resistance to heat-induced radiosensitization (39). Therefore, we determined the effects of NPM on DNA damage masking using this assay. In the halo assay DNA supercoil unwinding/rewinding is driven by increasing the concentration of the intercalating dye, propidium iodide. DNA strand breaks induced by ionizing radiation inhibit the ability of the loops to rewind supercoils in a dose dependent manner (40). The inability of DNA loops to be rewound is quantified as the difference between rewinding in the absence vs the presence of DNA damage (38,40). This difference referred to as the excess halo diameter (EHD) is estimated for PI concentrations between 10 and 50  $\mu$ g/ml (38). When cells are heated prior to irradiation, the EHD is reduced even though the amount of DNA damage per cell or per unit of DNA remains the same (4). Cells were untreated, irradiated with 10 Gy or heat shocked for 30 min at 45°C and then irradiated with 10 Gy prior to running the halo assay (Fig. 4A). Cells with normal NPM levels showed a typical heat-induced reduction in the inhibition of DNA supercoil rewinding following irradiation. In contrast, in cells with reduced NPM protein levels, hyperthermia had minimal effect on DNA supercoil rewinding after irradiation. Thus the masking of DNA damage, as detected by the halo assay, is reduced in cells with 80-90% lower NPM protein levels, suggesting that NPM plays a significant role in masking of damage when detected by the halo assay. The EHD (Fig. 4B, calculated as described above) for heated and irradiated cells with normal NPM levels was significantly reduced compared with radiation alone. In contrast, the heat-induced reduction of EHD, when NPM was knocked down, was ~ 80% less than that for normal NPM levels (Fig. 4B). Similar results were obtained

in HCC1806 cells (Fig. 4C). In summary a significant fraction (80%) of the masking effect, detected by the halo assay depends on the presence of NPM.

We used the ChIP assay to determine if NPM is associated with MAR, exon, or promoter DNA. NPM was immunoprecipitated with a NPM specific antibody from heated and control cells and the associated DNA analyzed by PCR using primers specific for MAR DNA, promoter DNA and exon DNA (31). Detection of specific DNA sequences (Fig. 5A) showed that NPM is associated with Topo I-MAR1, which increased more than 3-fold following hyperthermia (Fig. 5A & C). There appeared to be no association between NPM and the promoter DNA of c-jun or GAPDH, nor to Topo I exons 13 and 20 (Fig. 5C & D). Thus, only MAR DNA sequences were amplified by the PCR reactions on the DNA that co-precipitated with NPM, suggesting (albeit with limited sample size) that NPM binds MAR DNA, but not promoter or exon DNA. Further, the enhanced association between NPM and MAR DNA was reversed in a time dependent manner during post heat incubation (Fig. 5D), typical for post heat recovery from the masking effect (5). Since the ability to rewind DNA supercoils depends upon DNAnuclear matrix anchoring, the association of NPM with MAR DNA is consistent with the observation that the masking effect detected by the halo assay depends on NPM.

#### **DISCUSSION**

Here, we show that heat-induced NPM redistribution and DNA association contributes to the radiosensitization of cells, because multiple radiation sensitivity related endpoints (i.e., TER, inhibition of DSB repair and reduction of the EHD) are modulated in heated cells after NPM knockdown. The results also indicate that in heated cells NPM associates with MAR DNA, shown by modulation of the masking effect and the increased association of MAR DNA sequences with NPM. A model illustrating the effect of masking on DNA supercoil rewinding and the masking of DSBs is shown in Fig. 6. The hypothesized effects of DSB masking on downstream pathways and endpoints are illustrated in **Panel C**, assuming that if DNA damage is masked from a particular assay, then that damage will also be masked from (at least some of) the sensor steps in DNA repair. This assumption is supported by the observation that the duration of the initial DSB repair inhibition, 90 min, measured in the presence of PK corresponded to the duration of masking (reported previously, (6)) and the association of the TOPO I MAR with NPM.

It has been proposed that the nucleolus can act as repository of regulatory proteins that are released as needed in response to stress (17-19). Recent reports show that NPM binds chromatin in response to DNA double-strand breaks (41-42). These results suggest the importance of the nucleolus in modulating the cell's response to stress. The current work further supports the hypothesis that the nucleolus is a key element in the response to stress conditions by showing a role for NPM redistribution in heat-induced radiosensitization.

Since most clinical hyperthermia delivers 41-42°C temperatures, and NPM redistribution is induced by 45°C, the NPM radiosensitizing mechanism is not contributing to the TER achieved by current clinical hyperthermia. Therefore if the thermal effectiveness of a given heat exposure could be enhanced to induce NPM redistribution and DNA association, then a significant increase in TER would be expected. Thus, NPM redistribution and DNA association will be an excellent target for the development of thermal enhancing agents, reported previously (43-44).

In summary, after heat shock the nucleolar protein, NPM or B23 re-localizes from the nucleolus into the nucleoplasm increasing its association with MAR DNA, masks a fraction of X-ray induced DNA damage including DSB, inhibits their repair and contributes to heat-induced radiosensitization in two cell lines. This observation builds on previous reports that suggest

the nucleolus is a repository of stress responsive proteins. Thus NPM redistribution and MAR DNA binding should make a reasonable target for enhancing radiosensitization via thermal enhancers or mimetics.

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**(A)** Proteins were crosslinked to DNA by cisplatin from control and heat shocked HeLaS3 cells. Two forms of NPM, B23.1 and B23.2, are denoted by arrows. **(B)** HeLaS3 cells were incubated at 45°C for 45', fixed and stained with antibodies recognizing NPM. DNA was visualized with Hoechst dye and NPM with FITC filters. **(C)** Cells expressing a GFP-NPM fusion protein and counter stained with DAPI. **(D)** Redistribution of NPM following heat shock in HCC1806 cells.



#### **Figure 2. Reducing NPM reduces the TER**

**(A)** HeLaS3 cells were transfected with a NPM specific siRNA or a nonspecific siRNA, then irradiated with or without prior heat shock (30 min @ 45°C). The symbols are the average of 3-6 experiments  $\pm$  1 SD (n  $\leq$  3). The curves are normalized for cell killing by heat shock alone. **(B)** Heat shocked HeLaS3 cells were plated for cell survival and cell extracts prepared for protein analysis. **(C)** HeLaS3 cell extracts were probed for HSP70 and NPM. The average cell survival, after heat shock, is significantly lower for cells with lower NPM levels ( $p < 0.05$ , ttest). **(D)** Effect of NPM knockdown on the TER in the HeLaS3 cells and HCC1806 cells. See text for TER calculation.



#### **Figure 3. NPM contributes to DNA DSB repair inhibition and DSB masking**

**(A)** DNA DSBs were detected by the neutral comet assay. **Left Panel;** comet moment, measured after 30 min incubation with 0.27 U/ml PK. The initial damage was the same but a repair inhibition was evident in the heated sample. **Right Panel;** without PK digestion. the apparent reduction in initial damage shows that DNA DSBs are masked by a PK digestible component in heated cells (left arrow). Heat induced inhibition of repair can be seen as the increase in residual damage at 150 min. of repair (right arrow). **(B) Left Panel;** NPM knockdown reduces DNA DSB masking (right arrow) and repair inhibition (left arrow). The residual unrepaired DSBs are reduced by 50%. NPM knockdown had no effect on DNA DSB repair in the absence of heat shock. The **Right Panel**; the lack of effects for cells transfected

with a nonsense siRNA. **(C)** Masking of DNA DSB after 8 Gy is PK labile and partially reduced in cells with transient NPM knockdown as shown by reduction in comet moment. With versus without PK (p<0.05); NPM siRNA versus control or nonsense (NS) siRNA (p<0.05 experiments).



#### **Figure 4. Effect of NPM knockdown on DNA supercoil unwinding/rewinding after irradiation and/ or heat shock**

**(A)** The halo assay was used to determine the effects of NPM knockdown on the ability of nuclear DNA to undergo supercoiling changes after irradiation with or without prior heat shock. The points are a mean of 4-5 experiments. **(B)** Excess halo diameter (EHD) values were calculated from the rewinding portion of the 4-5 individual experiments and averaged (bars)  $\pm$  the SD. The only the non-transfected and the non-specific transfected showed a significant EHD reduction (p < 0.05, t-test) after heat shock. Inset: typical NPM levels. **(C)** EHD values for irradiated and heat plus irradiated HCC1806 cells. The non-transfected and the non-specific transfected cells showed a significant EHD reduction after heat shock.



#### **Figure 5. PCR reaction products from DNA associated with NPM**

**(A)** PCR results for the input DNA and that for the NPM-associated DNA recovered in the ChIP assay are indicated. Amplification of promoter sequences was detected in the input DNA but not in the DNA sequences co-immunoprecipitated with NPM. Control is marked "C"' heated at 45<sup>°</sup>C for 30 min is "O" and 30 min after heat is "30". NPM recovery is shown in **(B)**. Immunoprecipitates from unheated control cells are marked C. Immunoprecipitates from heat shocked (30 min at 45°C) cells are indicated by numbers, i.e.  $0 = 0$  min and 30 = 30 min after heat shock. After cells were heat shocked (45°C for 30 min) the CHIP assay was performed at the indicated time (**D**). Exon DNA was also analyzed. The relative change in NPM-Topo I MAR1 binding (open circles) and NPM-Topo I exon DNA (dotted line) is quantified (**C**).



**Figure 6. Model of the role of heat-induced NPM-DNA association in increased radiation sensitivity (A)** Illustration of the MAR region in an unheated cell with a radiation induced strand break that inhibits the rewinding of the right DNA loop. **(B)** Illustration of the MAR region in a heated cell with a radiation induced strand break that is buried in a protein-DNA aggregate and cannot inhibit the rewinding of the right DNA loop. **(C)** The downstream effects of the heat-induced masking of DNA damage which include inhibition of repair and increased radiation induced cell killing. Note that the 6-10 Kb range beyond the MAR is suggested by previous studies.