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Development and Validation of a Novel Assay to Identify Radiosensitizers that Target Nucleophosmin 1

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Abstract

A series of indole analogs that are synthesized using the scaffold of a potent radiosensitizer, YTR107, were tested for their ability to alter the solubility of phosphorylated nucleophosmin 1 (pNPM1). NPM1 is critical for DNA double strand break (DSB) repair. In response to formation of DNA DSBs, phosphorylated T199 NPM1 binds to ubiquitinated chromatin, in a RNF8/ RNF168-dependent manner, forming irradiation-induced foci (IRIF) that promote repair of DNA DSBs. A Western blot assay was developed using lead molecule, YTR107, for the purpose of screening newly synthesized molecules that target pNPM1 in irradiated cells. A colony formation assay was used to demonstrate the radiosensitization properties of the compounds. Compounds that enhanced the extractability of pNPM1 upon radiation treatment possessed radiosensitization properties.

Graphical Abstract

Novel Western blot assay was developed and tested for its applicability in identifying radiosensitizers that target nucleophosmin 1 (NPM1). Extractability of pNPM1 was used as a measurement for identification of active compounds from a library of analogs synthesized based on the lead molecule, YTR107.



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Introduction

Precision medicines that target specific driver mutations have revolutionized cancer therapy. Unfortunately, not all patients will be able to take advantage of these therapies. Many will present with tumors that do not express actionable molecular driver mutations. Non-small cell lung cancer (NSCLC) represents an excellent example, as 36% of patients fall into this category¹. Cytotoxic therapy continues to be a very important tool for the treatment of human cancers that do not express actionable molecular targets. Ionizing radiation is a cytotoxic agent that has a central role in cancer therapy and is used to provide local/regional control of cancer¹; a requirement for preventing tumor-mediated organ failure, tumor recurrence and metastasis^{2–4}. Recent advances in 3-D image-guided radiation therapy have significantly increased the probability of obtaining outstanding local/regional tumor control. However, a limitation to this therapy is the intrinsic radiation resistance of individual tumor cells⁵ due to increased DNA repair potential^{6–8}. Thus, targeting DNA repair represents a rational strategy for overcoming radiation resistance.

The indole structure of the radiation sensitizer, indomethacin⁹, was used as a scaffold for the synthesis of a series of (*Z*)-5-((*N*-benzyl-1*H*-indol-3-yl)methylene)imidazolidine-2,4-dione and (*Z*)-5-((*N*-benzyl-1*H*-indol-3-yl)methylene)pyrimidine-2,4,6(1*H*,3*H*,5*H*)-trione derivatives that incorporated a variety of aromatic substituents in both the indole and *N*-benzyl moieties. Functional phenotypic screening for structure activity relationships revealed that introduction of the electron withdrawing group 4-CN into the *N*-benzyl moiety yielded a potent radiosensitizing compound¹⁰, capable of sensitizing six NSCLC cell lines, HT29 colorectal cells, D54 glioblastoma cells, PANC1 pancreatic cancer cells, and two breast cancer cell lines^{11, 12}. This molecule, a substituted (*Z*)-5-((*N*-benzyl-1*H*-indol-3-yl)methylene)pyrimidine-2,4,6(1*H*,3*H*,5*H*)-trione was renamed YTR107 (Figure 1).

Inhibition of DNA double strand break (DSB) repair represented the mechanism responsible for YTR107-mediated radiosensitization¹². YTR107 exhibited efficacious radiosensitization in 2 tumor xenografts and a syngeneic tumor model but did not produce overt normal tissue toxicity¹¹ or normal tissue radiosensitization (unpublished results).

Use of YTR107 as a chemical probe resulted in the identification of the non-enzymatic chaperone, nucleophosmin 1 (NPM1) as a biological target that is critical for DNA DSB repair. In response to formation of DNA DSBs, phosphorylated T199 NPM1 binds to ubiquitinated chromatin, in a RNF8/RNF168-dependent manner, forming irradiation-induced foci (IRIF) that promote repair of DNA DSBs¹³. YTR107 binds to the amino terminus of NPM1, inhibiting IRIF formation, which in turn impairs DSB repair, and thus acts as a radiosensitizer. Genetic and cell biological approaches validated this concept by demonstrating that NPM1-deficient cells have impaired DNA DSB repair and consequently are radiosensitive. Use of NPM1-null mouse embryo fibroblasts demonstrated that the molecular basis of YTR107-mediated radiosensitization is YTR107 targeting of NPM1 and subsequent inhibition of DNA DSB repair. Although development of the YTR107 probe and discovery of its action represent a critical step for understanding a novel radiosensitizing mechanism, its use is hampered by limitations such as poor water-solubility.

In the present work, we report on a series of novel 2-methyl-*N*-benzyl aplysinopsin analogs, i.e. 2-methyl-5-((1-benzyl-1*H*-indol-3-yl)methylene)-2-oxodihydropyrimidine-4,6 (1*H*,5*H*)-triones and 2-methyl-5-((1-benzyl-1*H*-indol-3-yl)methylene)-2-thioxodihydro-pyrimidine-4,6(1*H*,5*H*)diones (Figure 2), which have been evaluated in a novel screening assay for their ability to modulate the extractability of phospho-nucleophosmin 1 (pNPM1 or pT199NPM1) after radiation treatment. The synthesis and anticancer properties of these analogues have recently been reported by us¹⁴. The screening assay was developed based on our novel observation that exposure of irradiated cancer cells to YTR107 increases the extractability of nuclear pNPM1 in high salt extraction buffer when compared to solvent control¹¹. In the present report, we analyzed the solubility of pNPM1 in NP-40 and RIPA

Methods and buffer compositions

these novel YTR107 analogs.

Lung cancer cells, A549, H460, and Calu1 as well as normal human lung fibroblasts, IMR-90 cells were purchased from ATCC and cultured in DMEM/F-12 (A549), RPMI-1640 (H460), and DMEM (Calu1 and IMR-90) media. Nuclei were isolated by suspending cells in nuclei isolation buffer on ice. Nuclear proteins were isolated by lysing the nuclei in high salt buffer. For total lysates, the cells were lysed either with NP-40 lysis buffer or RIPA buffer. The recipes to various buffers are given below.

buffers and correlated the findings with radiosensitization of H460 lung cancer cells with

NP-40 lysis buffer

50 mM Tris, pH 8.0, 150 mM NaCl, 0.01% NP-40, 0.1% SDS, 10 mM EDTA, 0.05% deoxycholic acid. Protease and phosphatase inhibitor cocktails were added before use.

RIPA buffer

10 mM Tris, pH 8.0, 150 mM NaCl, 5 mM EDTA, 0.1% SDS, 0.1% Triton X-100, 1% deoxycholic acid. Protease and phosphatase inhibitor cocktails were added before use.

High salt buffer

10 mM HEPES, pH 7.2, 420 mM NaCl, 0.1 mM EGTA, 1.5 mM MgCl2, 0.5 mM DTT, and 25% glycerol. Protease and phosphatase inhibitor cocktails were added before use.

Nuclei isolation buffer

50 mM Tris, pH 7.4, 50 mM NaCl, 10 mM EDTA, 0.5% NP-40. Protease and phosphatase inhibitor cocktails were added before use.

Western blot analysis

Lung cancer cells were grown to 70–80% confluency and were treated with analogs for 30 min, irradiated using Cs¹³⁷ source at 2 Gy per min and incubated 90 min in 37°C cell culture incubator. The cells were washed with cold PBS and protein was extracted with various buffers described above. Protein levels were measured using BioRad protein assay reagent (Cat# 500-0006). Equal amounts of protein were resolved on 10% SDS-PAGE and transferred onto a nitrocellulose membrane. The membrane was Western blotted for

nucleophosmin 1 and p-nucleophosmin 1 (T199) antibodies (pT199 NPM1 from Abcam and NPM1 from LifeTechnologies). Band intensities were quantified using Adobe Photoshop software.

Colony formation assay

This assay was performed as described in our previous publications^{12, 15}. In brief, cells were plated in T25 flasks, treated with drugs at 37° C for 30 min, irradiated and incubated for 90 min. The drugs were washed off with PBS and fresh media was added. The flasks were incubated 10–14 days. The colonies were stained with trypan blue and counted.

Extractability of pNPM1 in various buffers

In order to develop an assay to screen large number of molecules, it is necessary to demonstrate the simplicity of the assay with less processing steps. We have reported earlier on differential extractability of pNPM1 from irradiated cells with high salt buffer¹¹. Phosphorylation of NPM1 T199 is a highly specific cell cycle event that occurs just prior to initiation of S phase. In late G1, activated CDK2/cyclin E phosphorylates NPM1 at T199, causing NPM1 to dissociate from centrosomes, an event essential for centrosome duplication ¹⁶. CDK2/cyclin A activation during S and G2 insures continued phosphorylation of T199 NPM1¹⁷. During anaphase pNPM1 is dephosphorylated¹⁸, allowing it to associate with unduplicated centrosomes early in G1, and preventing centrosome duplication¹⁶. During the DNA damage response phosphorylation of T199 licenses NPM1 binding to ubiquitinated chromatin, in a RNF8/RNF168-dependent manner in response to formation of DNA DSBs¹³. Once bound to chromatin, pNPM1 forms irradiation-induced foci (IRIF) and promotes repair of DNA DSBs^{11–13}.

We have reported earlier that radiosensitizer, YTR107, enhances extractability of pNPM1 from nuclear extracts in high salt buffer upon radiation treatment¹¹. Dose dependent enhancement of pNPM1 solubility was determined for YTR107 using Western blots (Fig. 3). The band intensities were quantified using Adobe photoshop and EC₅₀ value was calculated from the graph generated from band intensities. In unirradiated A549 cells, pNPM1 can be extracted from the nuclei using a high salt buffer (data illustrating a representative immunoblot from A549 cells is shown in Figure 3, 0 Gy). However, pT199NPM1 cannot be extracted from irradiated cells by high salt buffer, as evidenced by a lack of immunoreactive protein (6 Gy, lane DMSO, Figure 3). This is a consequence of forming pNPM1 IRIF, as quantified by confocal microscopy¹². The confocal analysis confirmed that the change in buffer solubility upon irradiation was not a consequence of dephosphorylation of pNPM1¹². YTR107 enhanced the pNPM1 solubility at 10 μ M and above (Fig. 3). The EC₅₀ was determined to be 9.7 \pm 0.5 μ M. This experiment demonstrates that a Western blot assay can be used to identify possible radiosensitizers that target NPM1.

The nuclear extraction method is not suitable for the screening of large numbers of compounds, since it required isolation of nuclei, which is a laborious process and requires larger cell culture apparatus. In order to simplify the assay, we have tested two additional buffer compositions, NP-40 and RIPA buffers. This is a single step isolation of total cell lysates. Lead molecule, YTR107 was used to compare all three buffers in one assay to

demonstrate the identical outcome using Western blot (Fig. 4). Both, NP-40 and RIPA, buffers yielded similar results as that of nuclear extraction with high salt buffer.

Screening of New Compounds

Since we have demonstrated that the outcome from using various extraction buffers is comparable to each other, we have used NP-40 extraction method to screen newly synthesized YTR107 analogs (Western blot data not provided). NP-40 buffer is commonly used for protein isolation for Western blot analysis whereas RIPA buffer is routinely used for immunoprecipations. The band intensities were quantified using Adobe Photoshop and ratios were calculated between (Compound + 6 Gy) and (DMSO + 6 Gy). Ratios higher than 1 indicate that the molecule was capable of enhancing pNPM1 solubility upon radiation treatment (Table 1).

Dose response data obtained utilizing a molecule from the sub-library illustrated in Figure 2, PNR605 (**1n**), indicated that the EC₅₀ value to enhance extractability of pNPM1 is below 2.5 μ M (Fig. 5B). Identical extraction conditions were used for determining the EC₅₀ values for YTR107 and **1n**. Compound **1n** is thus 4 times more potent than the lead compound, YTR107, in enhancing the solubility of pNPM1 upon radiation. Similar observations were made when A549 cells were treated with PNR605 and total protein lysates were prepared with NP-40 lysis buffer (Fig. 5C).

In order to screen large number of compounds using the Western blot method, it is necessary to reduce the number of samples to be processed. So we have eliminated drug alone treatment and analysed the samples that were treated in combination with radiation. Solvent control (DMSO) was used to determine basal levels of pNPM1 present in the cancer cells (Figure 5). We have tested two concentrations of drug to evaluate the dose response of Calu1 cells. This approach also enables the determination of the dose-response of drugs in the extractability of pNPM1. For example, PNR584 (1c) at 2.5 μ M concentration was unable to alter the extractability when compared to DMSO + 6 Gy control (Figure 6). However, this molecule was able to alter pNPM1 solubility at 5 μ M. Similarly, PNR595 (1i) was unable to change the solubility of pNPM1 at both tested doses (Figure 6).

Normal cells will not respond to YTR107-mediated solubility of pNPM1

Recruitment of pNPM1 to chromatin is essential for the repair of radiation induced DSBs. During radiation treatment, normal tissue is also exposed to the radiation field, and may be sensitized by YTR107. We have used IMR90 normal lung fibroblasts to test whether YTR107 alters the extractability of pNPM1 upon radiation treatment. Western blot analysis revealed that YTR107 was unable to enhance the extractability of pNPM1 upon irradiation (Figure 7). This observation suggests that DSB repair is not hindered even when normal tissue is exposed to both drug and radiation. Thus, the extractability of pNPM1 seems to be specific to cancer cells.

Validation of Assay

In order to demonstrate that the compounds which increased the extractability of pNPM1 upon radiation treatment are radiosensitizers, we have randomly selected three compounds [PNR582 (**1b**), PNR605 (**1n**), and PNR596 (**1j**)] (Figure 2) and carried out a detailed analysis of radiosensitization effects on H460 cells. H460 cells were treated with the compounds and subjected to various doses of radiation. The cells were then subjected to colony formation assays (Figure 8). All three compounds are expected to be radiosensitizers. In fact, all three compounds enhanced the effect of radiation on H460 cells (Figure 8). Compound **1n** exhibited superior radiosensitization when compared to other two molecules.

Conclusion

The molecular chaperone nucleophosmin 1 (NPM1) is frequently overexpressed in cancer, where it supports oncogenesis-driven ribosome biogenesis¹⁹. Emerging research has suggests that NPM1 may be a critical factor for homologous recombination (HR)-mediated repair of DSBs. NPM1 is recruited to sites of DSBs, binds ubiquitinated chromatin and promotes RAD51-mediated HR¹³. Loss of HR imposes a radiosensitization phenotype on cells subjected to fractionated irradiation²⁰. Hence, we hypothesized that targeting pNPM1 would radiosensitize cancer cells. Since NPM1 is not an enzyme, screening of chemical libraries for NPM1 inhibitors would be challenging. We have observed that the extractability of pNPM1 is reduced when cancer cells are irradiated, and indole analog YTR107 reverses the effects of radiation on the extractability of pNPM1. Based on this observation, we have developed a Western blot assay to screen YTR107 analogs to identify molecules that could be effective radiosensitizers. The extractability of pNPM1 did not change with drug plus radiation treatment in normal lung fibroblasts, suggesting that the event is specific for cancer cells. Compounds identified through this assay radiosensitized lung cancer cells, suggesting the validity of the assay and the druggability of NPM1.

Acknowledgments

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Figure 1. Structures of YTR107 and its scaffold





1a-r

Compound	ID	R	X
1 a	PNR-5-81	Н	0
1b	PNR-5-82	Н	S
1c	PNR-5-84	4-CN	0
1d	PNR-5-85	4-CN	S
1e	PNR-5-87	4-COOCH3	0
1f	PNR-5-88	4-COOCH3	S
1g	PNR-5-90	3,4,5-trimethoxy	0
1h	PNR-5-91	3,4,5-trimethoxy	S
1i	PNR-5-95	3,5-dimethoxy	0
1j	PNR-5-96	3,5-dimethoxy	S
1k	PNR-6-01	2-Br	0
11	PNR-6-02	2-Br	S
1m	PNR-6-04	4-Br	0
1n	PNR-6-05	4-Br	S
10	PNR-6-07	2,4-dimethoxy	0
1p	PNR-6-08	2,4-dimethoxy	S
1q	PNR-6-10	2-CF ₃	0
1r	PNR-6-11	2-CF ₃	S

Figure 2.

Structures of 2-methyl-5-((1-benzyl-1*H*-indol-3-yl)methylene)-2-oxodihydropyri midine-4,6(1H,5H)-triones and 2-methyl-5-((1-benzyl-1*H*-indol-3-yl)methylene)-2-thioxodi hydropyrimidine-4,6(1H,5H)diones (**1a-r**).



Fig. 3.

YTR107 inhibits pNPM1 recruitment to chromatin in irradiated cells.

A Representative immunoblot illustrating the extraction of non-chromatin bound pT199NPM1 in 420 mM NaCl buffer from nuclei of A549 cells. Cells were exposed to DMSO or various concentrations of YTR107 for 30 min prior to, during, and for 90 min after 0 or 6 Gy. **B** Quantification of buffer soluble pNPM1 (\pm SD) from nuclei of irradiated cells relative to non-irradiated cells, N=3 experiments.



Figure 4. Effect of lysis buffers on the extractability of pNPM1 with YTR107

A549 cells were treated with drugs (25 μ M) for 30 min, irradiated, and incubated for 90 min. Nuclear extracts were made with high salt buffer (lanes 1–4). Total lysates using NP-40 buffer (lanes 5–8) and RIPA buffer (lane 9–12) were prepared and analysed by Western blot for pNPM1 and total NPM1.



Figure 5. Extractability of pNPM1 by compound 1n

H460 and A549 cells were treated with PNR605 (**1n**) for 30 min, irradiated (6 Gy), and incubated for 90 min. A) Nuclei were isolated from H460 cells and proteins were extracted with high salt buffer and immunoblotted. B) Bands were quantified from 5A using Adobe Photoshop and plotted using GraphPad Prism 4 software. EC_{50} was calculated at which 50% of pNPM1 extracted. C) Total lysates were prepared from A549 cells with NP-40 lysis buffer and immunoblotted for pNPM1 and NPM1 antibodies.

	2.5 5 2.5 5 2.5 5	2.5 5 2.5 5 2.5 5 2.5 5 μM	
			pNPM1
			GAPDH
DMSO DMSO+6 Gy	1b + 6 Gy 1b + 6 Gy 1c + 6 Gy 1c + 6 Gy 1d + 6 Gy 1d + 6 Gy	$\begin{array}{c} 1e+6 \ Gy \\ 1e+6 \ Gy \\ 1f+6 \ Gy \\ 1f+6 \ Gy \\ 1i+6 \ Gy \\ 1j+6 \ Gy \\ 1j+6 \ Gy \\ 1j+6 \ Gy \end{array}$	

Figure 6. Extractability of pNPM1 in Calu1 cells after irradiation in the presence of various YTR107 analogs

Calu1 cells were treated with DMSO/drugs for 30 min, irradiated, and incubated for 90 min. Total proteins were isolated with NP-40 lysis buffer and analyzed by Western blots.



Figure 7. Effect of YTR107 on IMR90 cells

IMR90 cells were treated with YTR107 (25 μ M) for 30 min, irradiated, and incubated for 90 min. Total lysate was prepared with NP-40 lysis buffer and analyzed by Western blot.



Figure 8. Radiosensitization of H460 cells by YTR107 analogs

H460 cells were treated with 5 μ M of 1n, 1b, or 1j for 30 min, irradiated, and incubated for 90 min. The drugs were then washed off and the flasks were incubated for 10–14 days, stained with trypan blue and colonies were counted and analyzed.

Table 1

Band intensities were quantified from Western blots and ratio of band intensity from radiation + drug was divided by DMSO + radiation. Higher values indicate that the extractability pNPM1 is enhanced upon treatment with drug.

Compound	Ratio in A549 Cells (Compound+6Gy/DMSO+6Gy) 25 µM	
DMSO	1.00	
1a	1.53	
1b	1.96	
1c	1.58	
1d	1.86	
1e	1.48	
1f	1.74	
1g	1.16	
1h	1.97	
1i	1.63	
1j	1.97	
1k	1.05	
11	1.68	
1m	1.23	
1n	1.78 with 2.5 µM drug	
10	ND [#]	
1p	(1.22)	
1q	1.16	
1r	1.54	
YTR107	1.51	

[#]ND – Not determined