## Supplemental Data

Cisplatin in Combination with MDM2 Inhibition Downregulates Rad51 Recombinase in a Bimodal Manner to Inhibit Homologous Recombination and Augment Tumor Cell Kill

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Supplemental Table S1. Statistical analysis by post hoc Tukey of Rad51 mRNA expression following exposure of A2780 cells to cisplatin, Nultin-3 or the combination

| Control vs. Cisplatin | Diff $=-0.0543$ | $95 \% \mathrm{Cl}=-0.2426$ to 0.1341 | $p=0.7939$ |
| :--- | :--- | :--- | :--- |
| Control vs. Nutlin-3 | Diff $=-0.2853$ | $95 \% \mathrm{Cl}=-0.4736$ to -0.0969 | $p=0.0056$ |
| Control vs. Combination | Diff $=-0.6130$ | $95 \% \mathrm{Cl}=-0.8014$ to -0.4246, | $p=0.0000$ |
| Cisplatin vs. Nutlin-3 | Diff $=-0.2310$ | $95 \% \mathrm{Cl}=-0.4193$ to -0.0426, | $p=0.0184$ |
| Cisplatin vs. Combination | Diff $=-0.5587$ | $95 \% \mathrm{Cl}=-0.7471$ to -0.3704 | $p=0.0000$ |
| Nutlin-3 vs. Combination | Diff $=-0.3278$ | $95 \% \mathrm{Cl}=-0.5161$ to -0.1394 | $p=0.0023$ |

Results were subjected to one-way ANOVA and then Tukey post hoc. Cl , confidence interval.

Supplemental Table S2. Statistical analysis by post hoc Tukey of Rad51 mRNA expression following exposure of A2780 p53-proficient (+/+) cells to cisplatin, Nultin-3 or the combination

| Control vs. Cisplatin | Diff $=-0.0571$ | $95 \% \mathrm{Cl}=-0.2891$ to 0.1749 | $\mathrm{p}=0.8578$ |
| :--- | :--- | :--- | :--- |
| Control vs. Nutlin-3 | Diff $=-0.2601$ | $95 \% \mathrm{Cl}=-0.4921$ to -0.0281 | $\mathrm{p}=0.0291$ |
| Control vs. Combination | Diff $=-0.6163$ | $95 \% \mathrm{Cl}=-0.8483$ to -0.3842 | $\mathrm{p}=0.0001$ |
| Cisplatin vs. Nutlin-3 | Diff $=-0.2030$ | $95 \% \mathrm{Cl}=-0.4350$ to 0.0290 | $\mathrm{p}=0.0880$ |
| Cisplatin vs. Combination | Diff $=-0.5591$ | $95 \% \mathrm{Cl}=-0.7912$ to -0.3271 | $\mathrm{p}=0.0002$ |
| Nutlin-3 vs. Combination | Diff $=-0.3561$ | $95 \% \mathrm{Cl}=-0.5881$ to -0.1241 | $\mathrm{p}=0.0051$ |

Results were subjected to one-way ANOVA and then Tukey post hoc. CI, confidence interval.

## Supplemental

Figure S1


B


2780CP/Cl-16

Supplemental Figure S1 Legend. (A) A2780 cells were treated with cisplatin, Nutlin-3 or the combination for 24 hours, collected and analyzed by immunoblot with the indicated antibody. Drug concentrations were as follows: $2780 \mathrm{CP} / \mathrm{Cl}-16$ (cisplatin, $5 \mu \mathrm{M}$; Nutlin-3, $2 \mu \mathrm{M}$ ) and $2780 \mathrm{CP} / \mathrm{Cl}-24$ (cisplatin, $5 \mu \mathrm{M}$; Nutlin-3, $4 \mu \mathrm{M}$ ). (B) $2780 \mathrm{CP} / \mathrm{Cl}-16$ cells stably transfected with DNA damage reporter plasmid were treated with cisplatin, Nutlin-3 or combination for 24 hours, cells were then fixed and DNA damage foci monitored using confocal microscopy. (C) $2780 \mathrm{CP} / \mathrm{Cl}-24$ were treated as in Fig. S1B to monitor DNA damage foci.

Fig. S2
A




B


C


D


Supplemental Figure S2 Legend. (A) A2780, 2780CP/Cl-16 and 2780CP/Cl-24 cells were treated with cisplatin $(200 \mu \mathrm{M})$ for 2 hours or pretreated with Nutlin-3 for 24 hours following 2 hours cisplatin $(200 \mu \mathrm{M})$, cells were then collected and intracellular cisplatin concentration determined by atomic absorption. The results are shown as mean $\pm$ SD of three independent experiments. (B) A2780 cells treated for 24 hours with cisplatin $(1 \mu \mathrm{M})$, Nutlin- $3(0.5 \mu \mathrm{M})$ or the combination were subjected to RPPA analysis, and the proteins associated with DNA repair pathways and identified by Ingenuity Pathway Analysis are shown as fold change vs. control. (C) U2OS cells were treated with cisplatin $(7.5 \mu \mathrm{M})$, Nutlin- $3(0.5 \mu \mathrm{M})$ or combination for 24 hours, cells were then harvested and analyzed using immunoblot with indicated antibodies. (D) Rad51 levels in A2780 (cisplatin, $1 \mu \mathrm{M}$ and Nutlin-3, $0.2 \mu \mathrm{M}$ ), 2780CP/Cl-16 (cisplatin, $5 \mu \mathrm{M}$ and Nutlin-3, $0.5 \mu \mathrm{M}$ ) and $2780 \mathrm{CP} / \mathrm{Cl}-24$ (cisplatin, $5 \mu \mathrm{M}$ and Nutlin-3, $0.5 \mu \mathrm{M}$ ) were determined by Western blot following drug exposure for 24 hr . (E) Correlation between MDM2-pS166 and RAD51 from TCPA database using information from the category 'MDACC ovary level 4 cell lines.' The values shown are expressed as Log 2.

