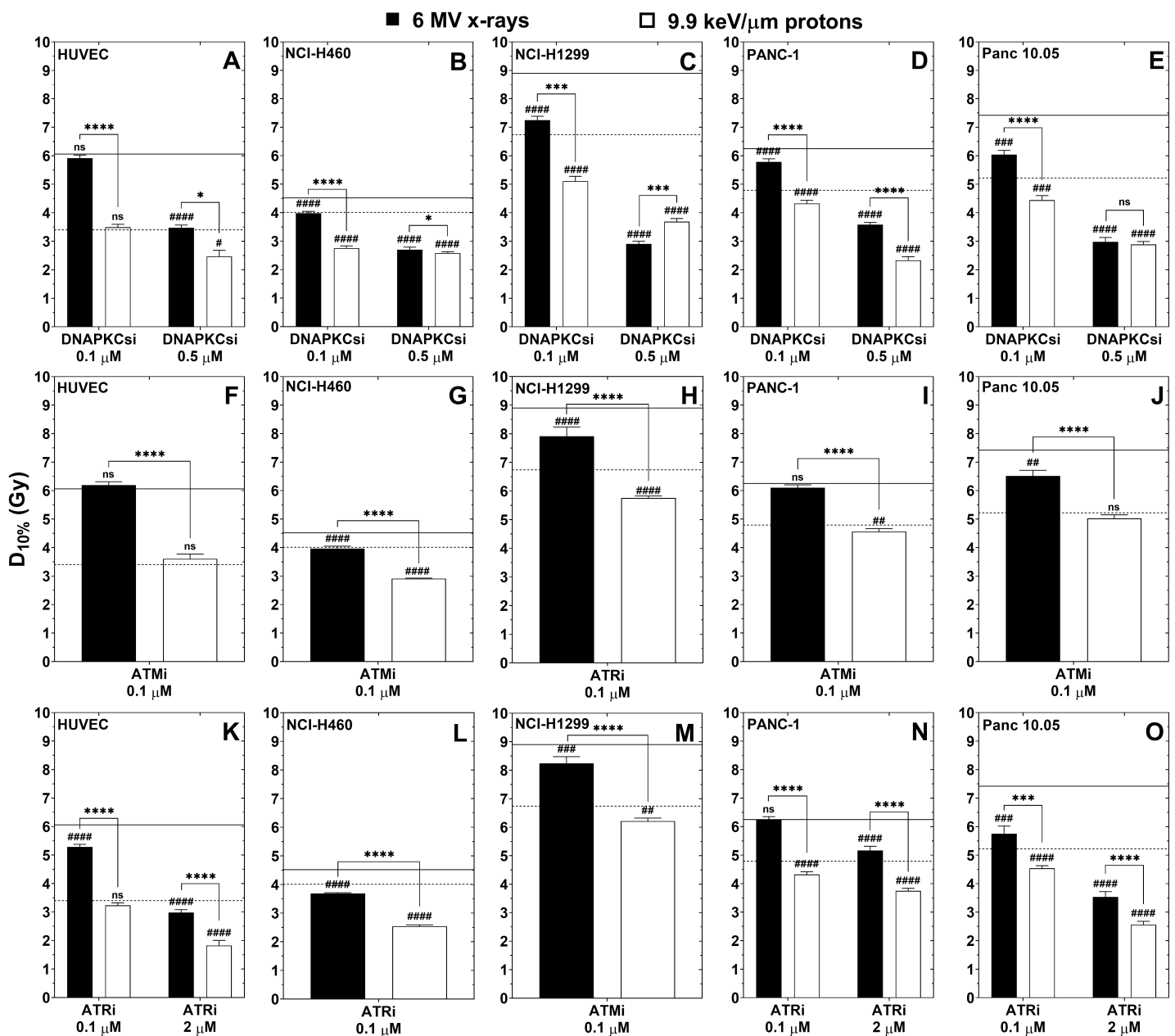
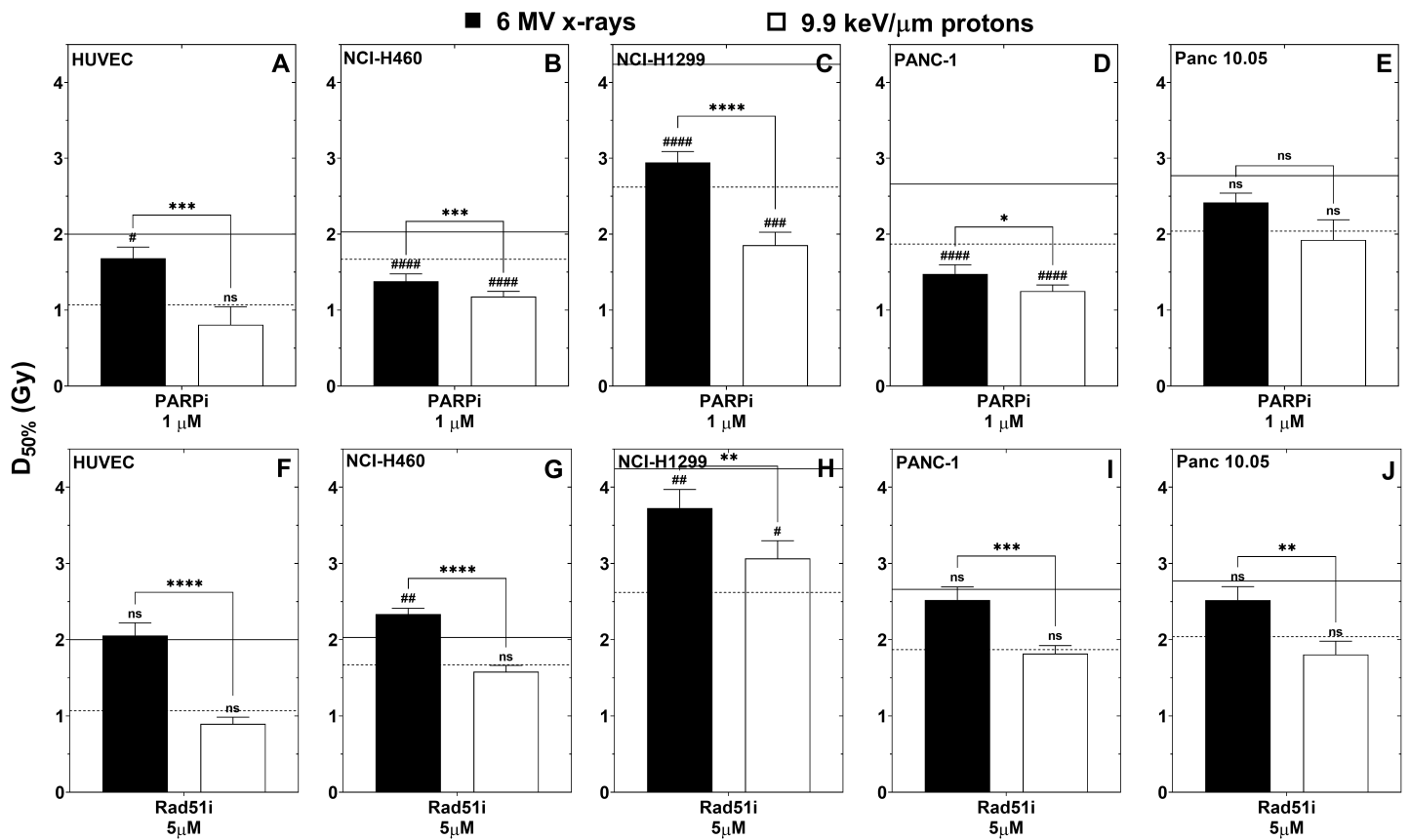


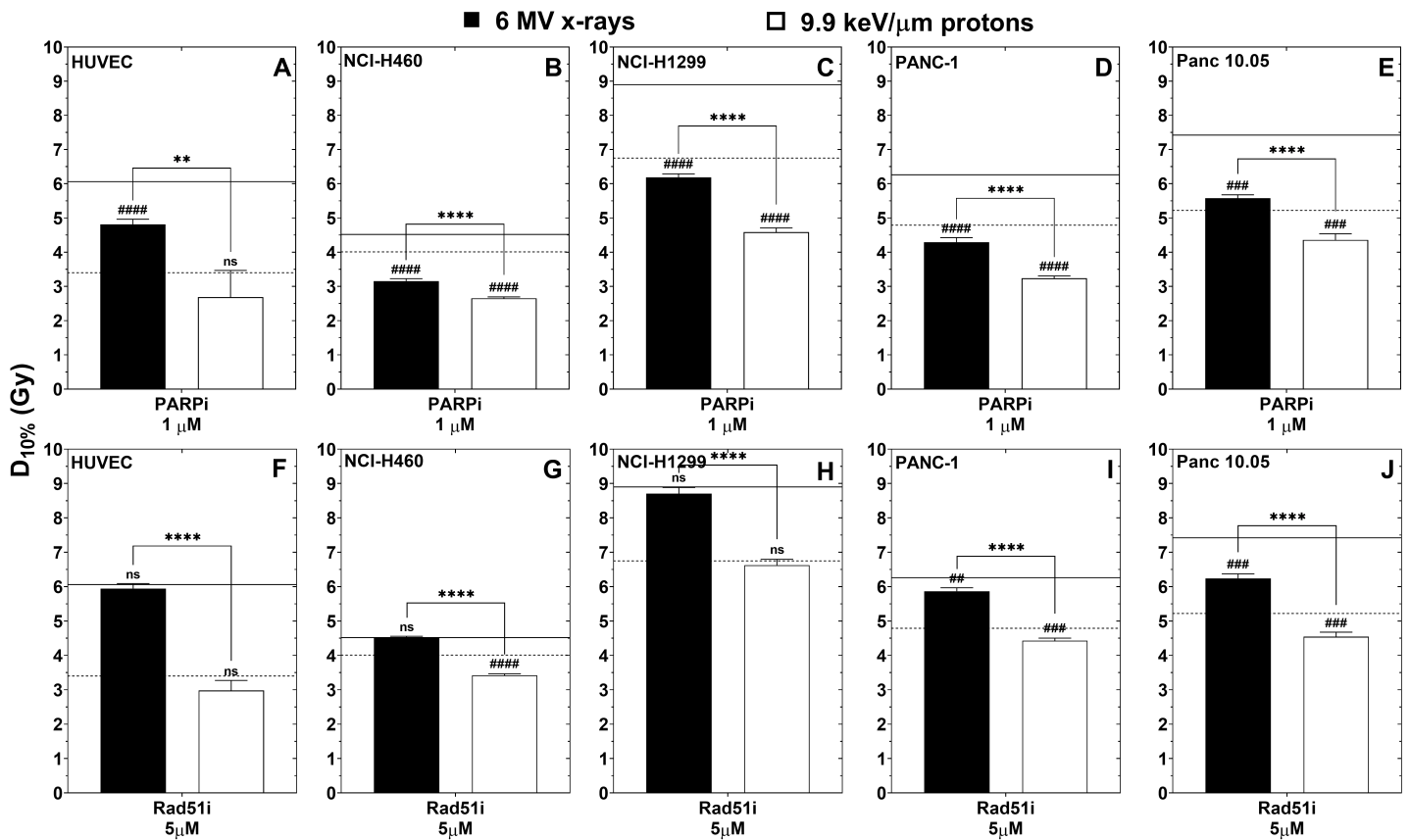
Supplementary Figure 1:  $D_{50\%}$  for combination of DNA repair inhibitors with x-rays or protons. (A-D) RT+DNAPKcsi; (F-J) RT+ATMi; and (K-O) RT+ATRi for HUVEC, NCI-H460, NCI-H1299, PANC-1 and Panc 10.05 cell lines. Lines represent responses to x-rays (solid) and protons (dashed) when treated with DMSO. Number (N) of biological repeats for each condition is given in Figure 2. Significance denotes differences between x-rays and protons with a given inhibitor (\*) or differences between a given inhibitor/radiation pair and its respective radiation type with DMSO (respectively (#). Error bars represent the standard deviation. ns: non-significant; \* or #:  $p < 0.05$ ; \*\* or ##:  $p < 0.01$ ; \*\*\* or ###:  $p < 0.001$ ; and \*\*\*\* or ####:  $p < 0.0001$ .



Supplementary Figure 2: D<sub>10%</sub> for combination of DNA repair inhibitors with x-rays or protons. (A-D) RT+DNAPKcsi; (F-J) RT+ATMi; and (K-O) RT+ATRi for HUVEC, NCI-H460, NCI-H1299, PANC-1 and Panc 10.05 cell lines. Lines represent responses to x-rays (solid) and protons (dashed) when treated with DMSO. Number (N) of biological repeats for each condition is given in Figure 2. Significance denotes differences between x-rays and protons with a given inhibitor (\*) or differences between a given inhibitor/radiation pair and its respective radiation type with DMSO (respective line) (#). Error bars represent the standard deviation. ns: non-significant; \* or #: p<0.05; \*\* or ##: p<0.01; \*\*\* or ###: p<0.001; and \*\*\*\* or ####: p<0.0001.



Supplementary Figure 3: D<sub>50%</sub> for combination of DNA repair inhibitors with x-rays or protons. (A-D) RT+PARPi; and (F-J) RT+Rad51i for HUVEC, NCI-H460, NCI-H1299, PANC-1 and Panc 10.05 cell lines. Lines represent responses to x-rays (solid) and protons (dashed) when treated with DMSO. Number (N) of biological repeats for each condition is given in Figure 3. Significance denotes differences between x-rays and protons with a given inhibitor (\*) or differences between a given inhibitor/radiation pair and its respective radiation type with DMSO (respective dotted line) (#). Error bars represent the standard deviation. ns: non-significant; \* or #: p<0.05; \*\* or ##: p<0.01; \*\*\* or ###: p<0.001; and \*\*\*\* or ####: p<0.0001.



Supplementary Figure 4:  $D_{10\%}$  for combination of DNA repair inhibitors with x-rays or protons. (A-D) RT+PARPi; and (F-J) RT+Rad51i for HUVEC, NCI-H460, NCI-H1299, PANC-1 and Panc 10.05 cell lines. Lines represent responses to x-rays (solid) and protons (dashed) when treated with DMSO. Number (N) of biological repeats for each condition is given in Figure 3. Significance denotes differences between x-rays and protons with a given inhibitor (\*) or differences between a given inhibitor/radiation pair and its respective radiation type with DMSO (respective dotted line) (#). Error bars represent the standard deviation. ns: non-significant; \* or #:  $p < 0.05$ ; \*\* or ##:  $p < 0.01$ ; \*\*\* or ###:  $p < 0.001$ ; and \*\*\*\* or ####:  $p < 0.0001$ .