

Comparison of phosphorylation kinetics in DNA repair proteins after exposure to high and low LET radiations

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We irradiated plateau phase normal human fibroblasts with 2 Gy X-rays, 70 keV/um carbon (290MeV/n) and 200 keV/um iron ions (500 MeV/n) and observed the kinetics of phosphorylation in various proteins associated with DNA double strand break (DSB) repair. GammaH2AX foci, a marker for DSBs, were detected immediately after irradiation, and the peak of phosphorylation was seen 30 to 60 min post-irradiation for three kinds of radiations. Disappearance of gamma-H2AX foci was much faster for X-irradiated samples than that for heavy ion irradiated samples; the phosphorylation kinetics for carbon and iron ions are similar for gamma-H2AX foci. In contrast, phosphorylation of an NHEJ protein, DNA-PKcs (threonine 2609) was significantly delayed in carbon and iron irradiated cells when compared to X-irradiated cells. Disappearance of DNA-PKcs sites was much faster in X-irradiated samples than carbon and iron samples which showed a similar pattern as in the case of gamma-H2AX. Furthermore, in the case of ATM protein phosphorylation (serine 1981), iron irradiation alone caused a significant initial delay, but the kinetics of disappearance is similar for iron and carbon samples with much higher remaining number of foci in iron samples than those for X-rays and carbon ions.

These results suggest that 1) high LET irradiation induces complex and/or severe DNA DSB damage which affects the function of DSB repair proteins, 2) Both ATM and DNA-PKcs may recognize the complexity of DSBs, but ATM may be more sensitive to detecting the complexity of DSB damage, 3) gamma-H2AX may just recognize DSBs, but not the complexity of them. Our studies also indicate that high LET radiation would be a useful tool to elucidate the complexity of initial DSB damage and the mechanism of repair on these damages.