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Differences in DNA damage and repair in human cancer and normal cells after ultrashort pulsed electron beam irradiation

Targeting the DNA Damage Response in CML cells

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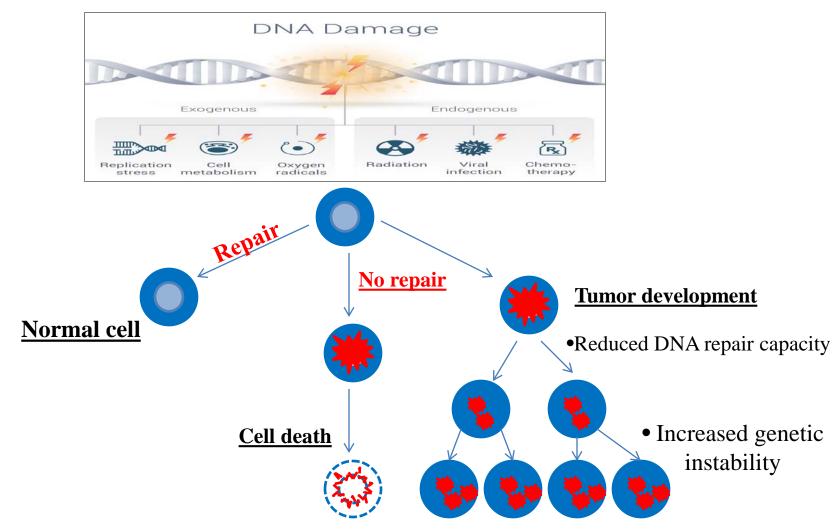




Research overview:

DNA damage response in cancer cells

Every cell experiences up to 10⁵ spontaneous or induced DNA lesions per day

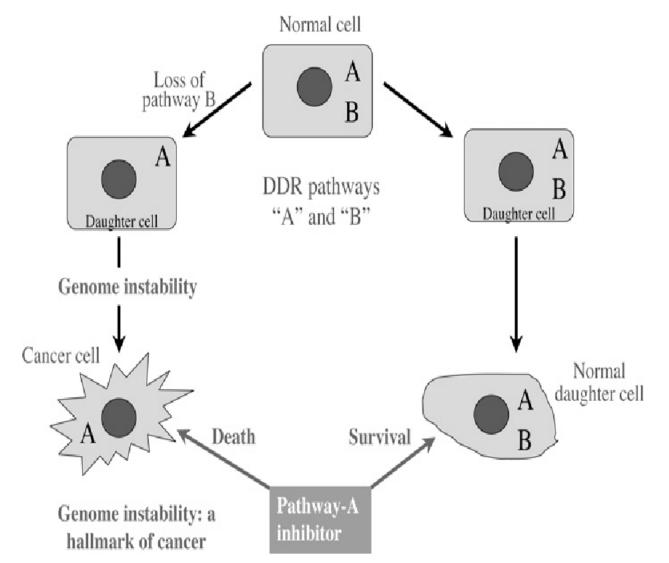




Targeting DNA damage response



in cancer therapy



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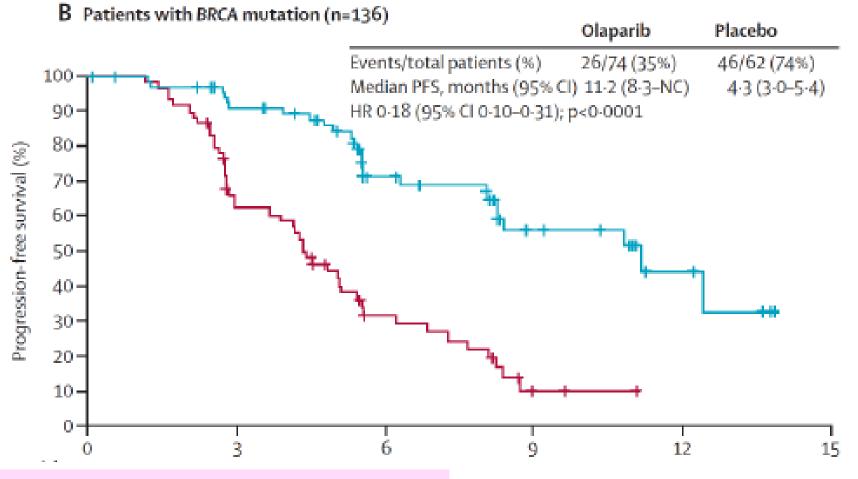


<u>Synthetic lethal interactions</u> in DNA repair genes implicated in cancer

Pathway	Protein	Syndrome	Primary cancers			
HR	BRCA1		breast, ovarian			
	BRCA2	Fanconi's anemia	breast, ovarian			
	AD54B		non-Hodgkin lymphoma, colon cancer			
NHEJ	(EJ					
	MRE11	Ataxia-telangiectasia- like disorder	colorectal cancer			
	LIG4	LIG4 syndrome	Leukemia			
	Artemis	Omenn syndrome	Lymphoma			
NER						
	XPA	Xeroderma pigmentosum	Skin cancers			
	ERCC1	cerebro-oculo-facio-skeletal syndrome	squamous cell carcinoma, head /neck			
Crosslink repair						
	FANCA, B, C, D2, E	Fanconi's anemia	Various			
etc.						

© Helleday et al., Nature Reviews Cancer, 2008, vol 8, no. 3, pp. 193-204. DOI: 10.1038/nrc2342





EMA approval: 16th December 2014

FDA approval: 19th December 2014

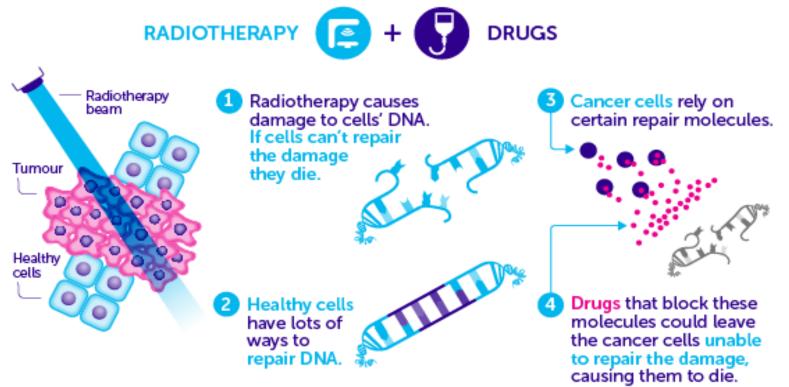




Targeting DDR in cancer radiotherapy

COMBINING RADIOTHERAPY WITH DRUGS

Drugs that stop cancer cells repairing their DNA could help make radiotherapy more effective.



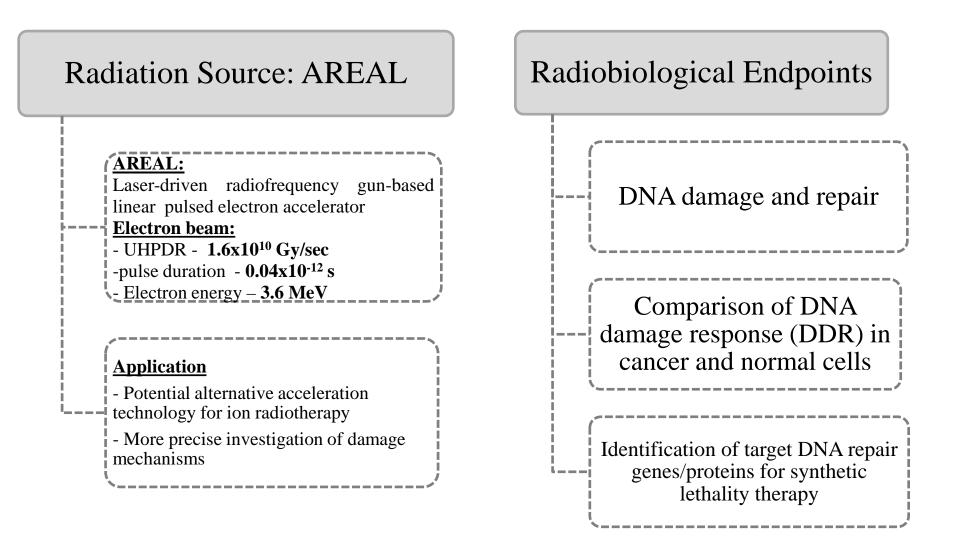


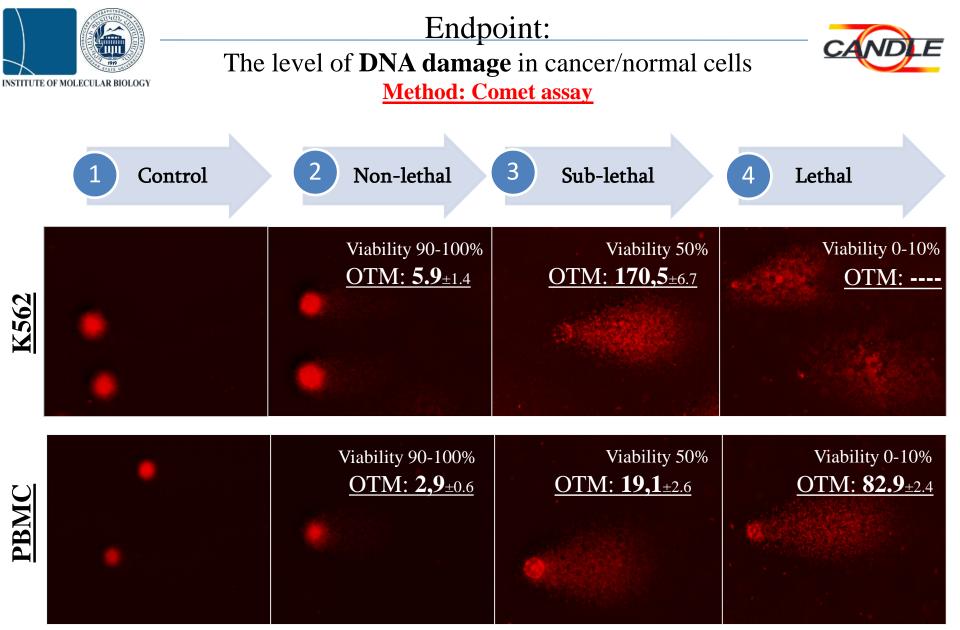




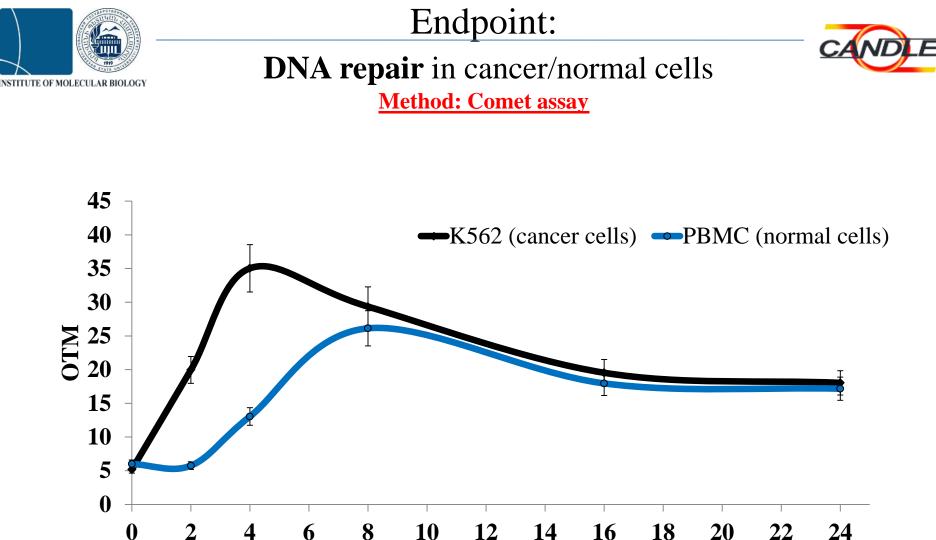
The Aim:

DDR in cancer and normal cells after UPEB radiation





Level of DNA damage in human normal (PBMCs, blood normal cells, female) and cancer (K562, blood cancer cells, female) cells after irradiation (0 h) at the non-lethal, sub-lethal and lethal doses **OTM** - Olive Tail moment is defined as the product of the tail length and the fraction of total DNA in the tail



Dose, Gy

The level of DNA-damage in PBMC (normal) and K562 cell line (cancer) cells **after 3 hours of irradiation**

OTM - Olive Tail moment is defined as the product of the tail length and the fraction of total DNA in the tail

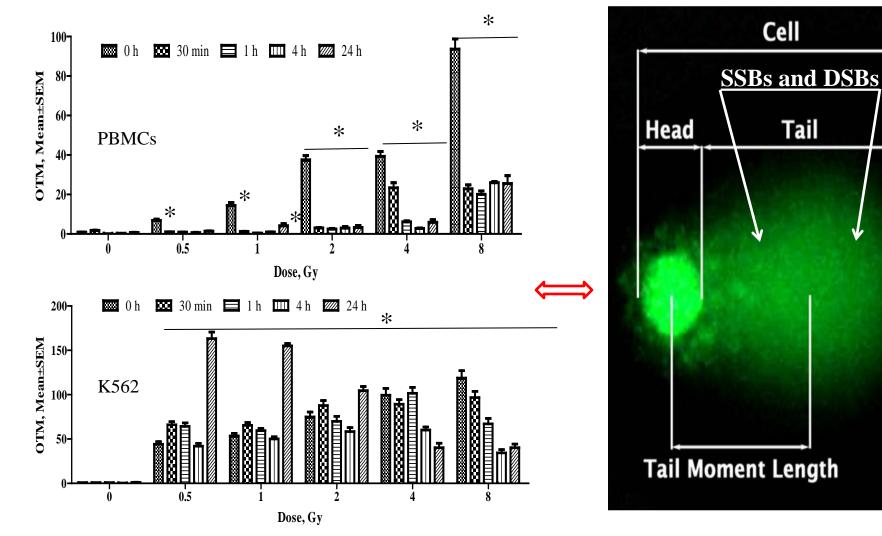


Endpoint:

DNA repair kinetics in cancer/normal cells



Method: Comet assay

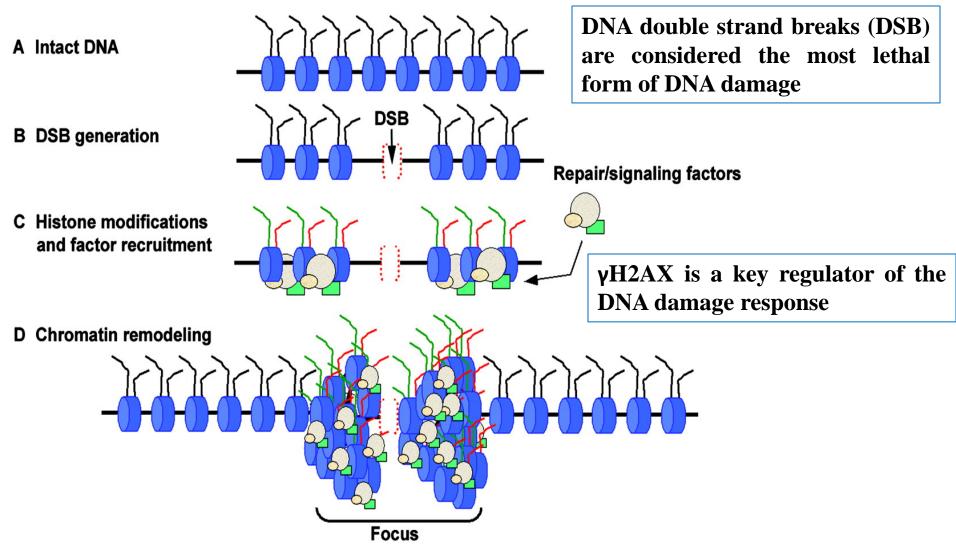


Comparison of the induced primary DNA-damage level and repair kinetics in PBMCs (a) and K562 (b) cells. *p<0.05 in comparison to corresponding control





Phosphorylation of Histone H2AX at DNA Double-Strand Breaks



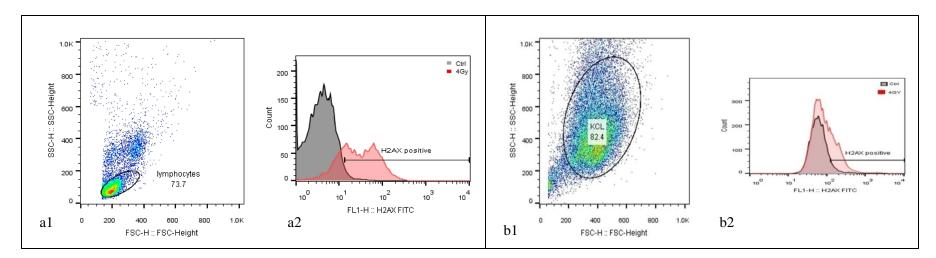


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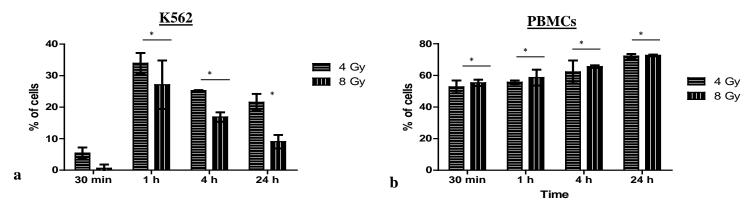


DNA DSBs and repair kinetics in cancer and normal cells

Method: Flow cytometric analysis of γ -H2AX



The representative dot-plot (a1, b1) and histogram (a2, b2) of the level of γ -H2AX in PBMCs (a) and K562 cells (b) after sub-lethal (4 Gy) dose of irradiation



The kinetics of γ -H2AX foci formation in cancer (a) and normal (b) cells after lethal (8 Gy) and sub-lethal (4 Gy) doses of irradiation. *- p<0.05 in comparison with non-irradiated cells





Ionizing radiation induced-DNA SSBs/DSBs repair pathways

Non-Homologous End Joining <u>NHEJ</u>





Homologous Recombination Repair <u>HRR</u>

Slow <u>Free of errors</u> Core protein: MRE11

Base Excision Repair
BER

FAST <u>Free of errors</u> Core protein: APEX1

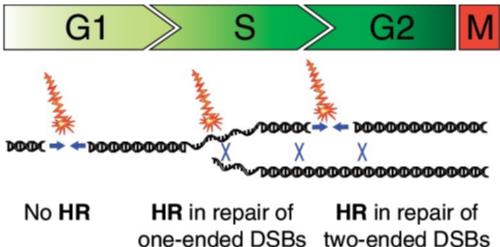


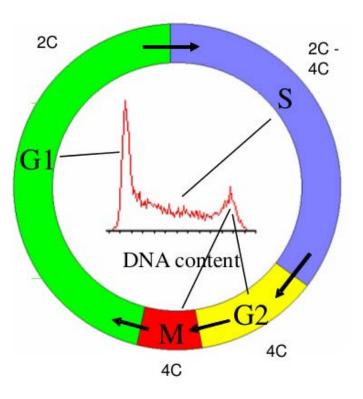
DNA SSBs/DSBs repair pathway activation accompanied with

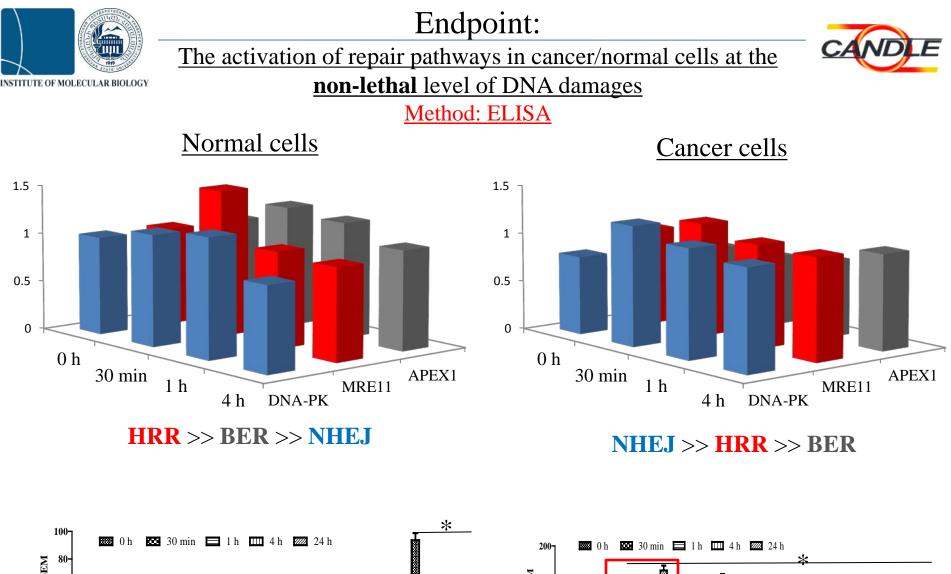
cell cycle arrest

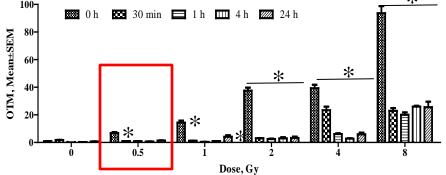


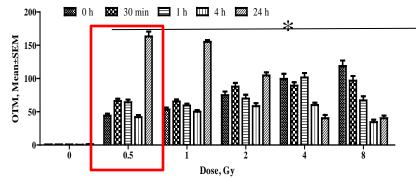
- G1 Growth
- S DNA synthesis
- G2 Growth and preparation for mitosis
- M Mitosis (cell division)

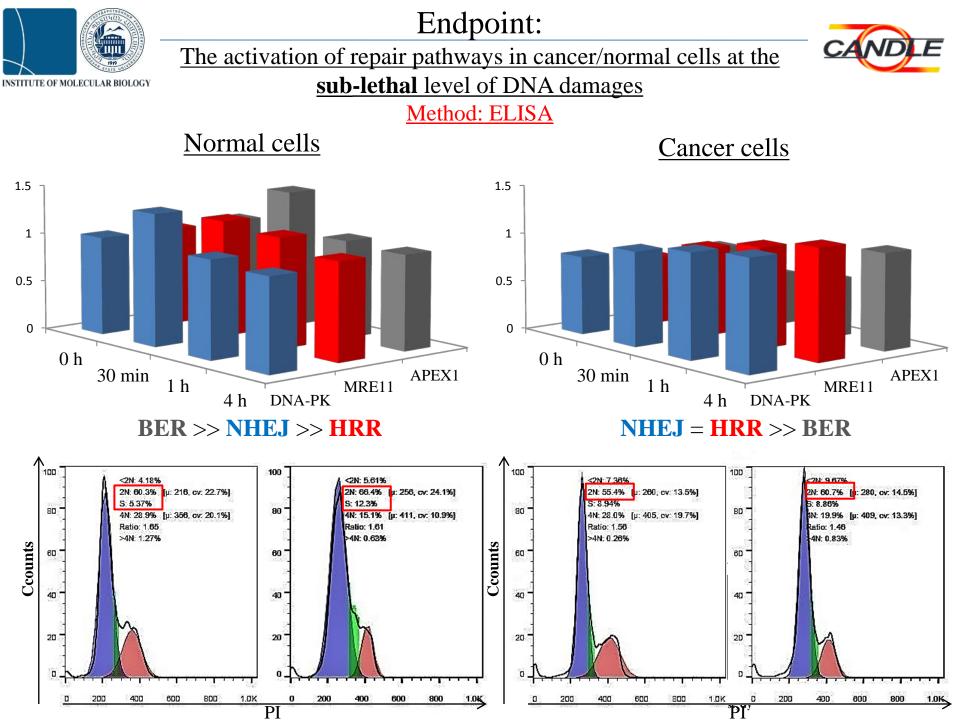


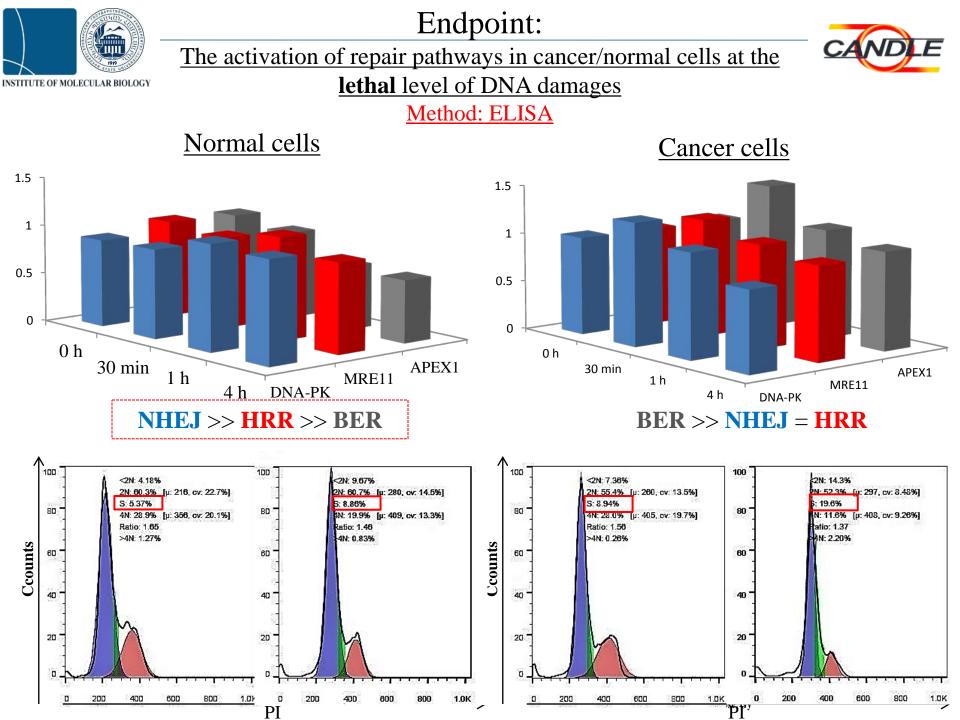










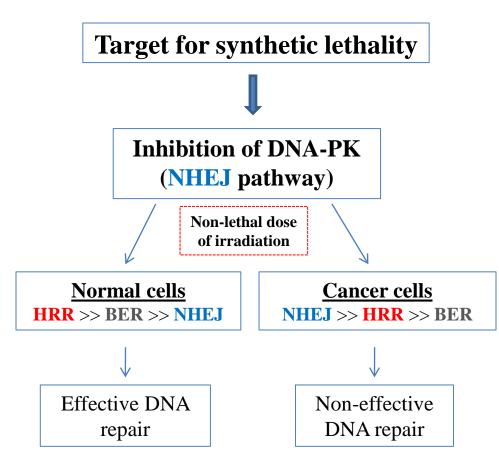


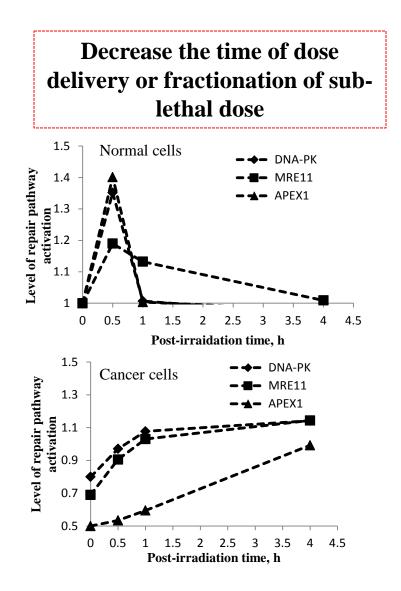


Conclusions/Recommendations

<u>To increase the sensitivity of</u> <u>cancer cells (CML) vs. normal cells</u>







Acknowledgements







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Thanks for Your Attention !!!