

# A Systematic Review of the Proton and Carbon FLASH Effect

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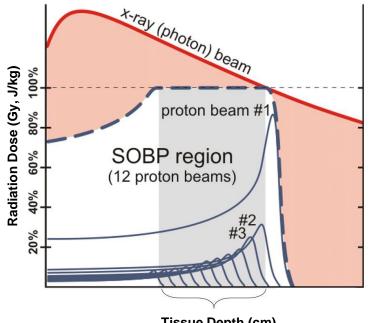
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### **Tissue Depth (cm)**

## **Radiation Therapy**

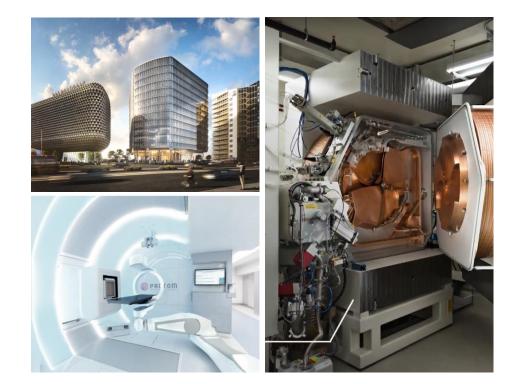
- RT most common cancer treatment, ~50% patients in developed countries
- Limiting factor healthy tissue damage
- Particle therapy benefits:
  - Protect more healthy tissues
  - Reduces damage to critical structures
  - Good for radioresistant tumours
  - Paediatrics many years to manifest symptoms, emphasis on reducing normal tissue damage

Image courtesy of the South Australian Medical Heritage Society



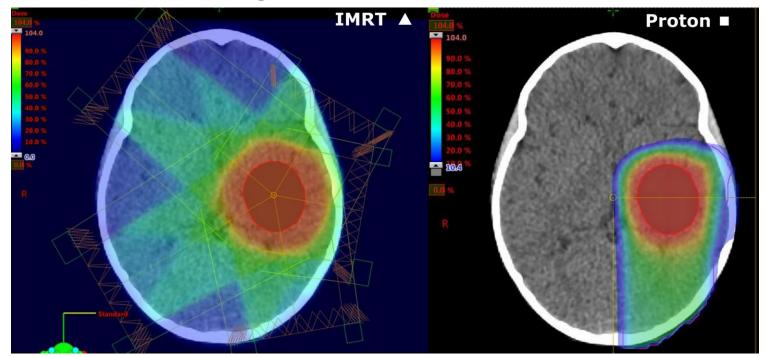
## **Australia First!**

- Australian Bragg Centre for Proton Therapy (SAHMRI 2)
- Expected completion: 2023
- First patient treatments: 2024
- Will contain ProTom's Radiance 330 akin to Massachusetts General Hospital – synchrotron
- Currently: 16.5 MeV GE PETtrace cyclotron
- Will be using for my experiments





## **X-rays versus Protons**



UniSA STEM

Image courtesy of Mikaela Dell'Oro and Eva Bezak

## **X-rays versus Protons**

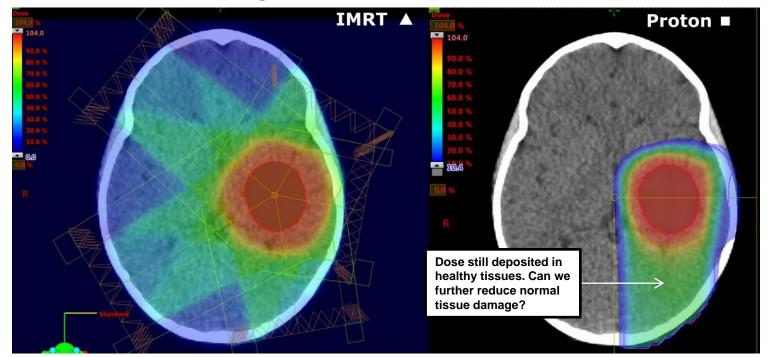


Image courtesy of Mikaela Dell'Oro and Eva Bezak

## What is FLASH?

Use of single pulses of radiation at ultra-high dose rates to treat solid tumours (>40 Gy/second)

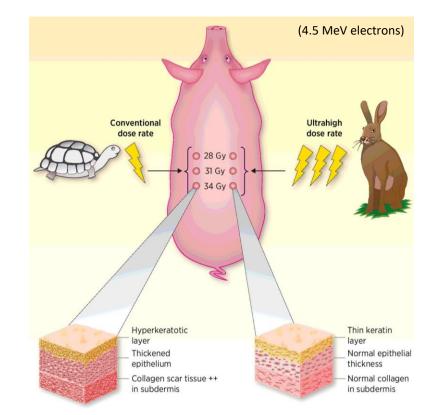
Contrasts conventional (CONV) therapy which uses low dose rates (Gray/minute) in multiple pulses or 'fractions'

Equivalent tumour kill efficacy, but improves the sparing of healthy, normal tissues

Biological and physical mechanisms unclear...

### **Predicted mechanisms:**

- 1. Oxygen depletion
- 2. DNA Damage Repair & Response
- 3. Immune Cell Modulation and Inflammation



Vozenin MC et al. 2019, Clin Cancer Res



- What is it?
  - A reproducible method to collect all available research on a specific topic

dentification

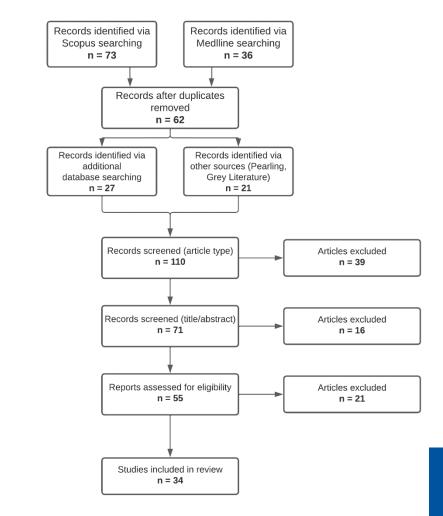
Eligibility

and

Screening

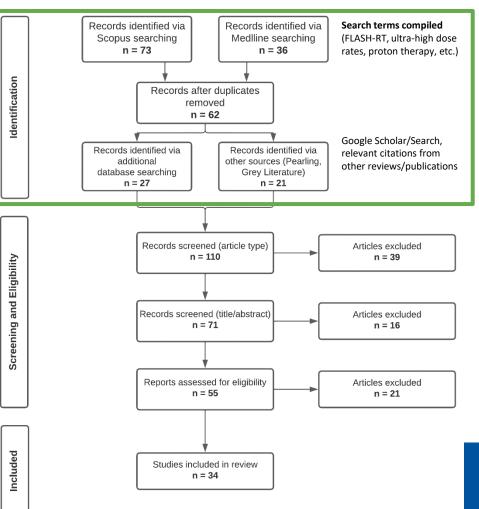
Included

- Review follows PRISMA Guidelines
- Preferred Reporting Items for Systematic Reviews and Meta Analyses
- Titled: The Current Status of FLASH Particle Therapy: A Systematic Review
- Exclusion criteria:
  - 1. Only proton and/or carbon FLASH papers
  - 2. Must cite use of > 40 Gy/s
  - 3. Radiobiological effect measured



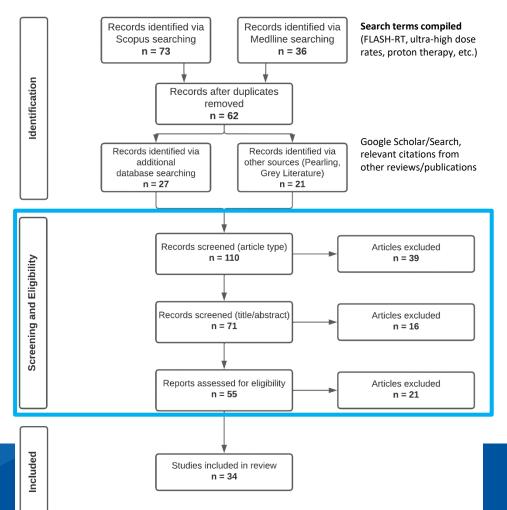


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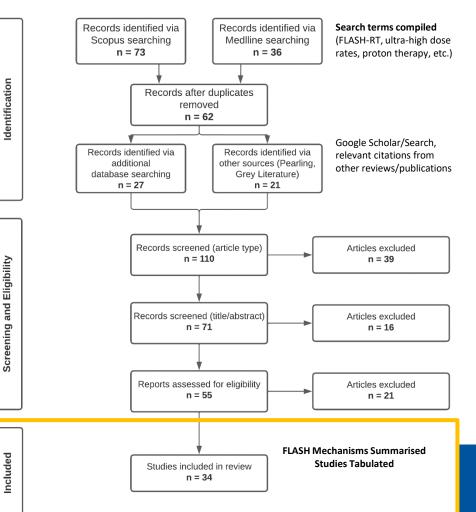


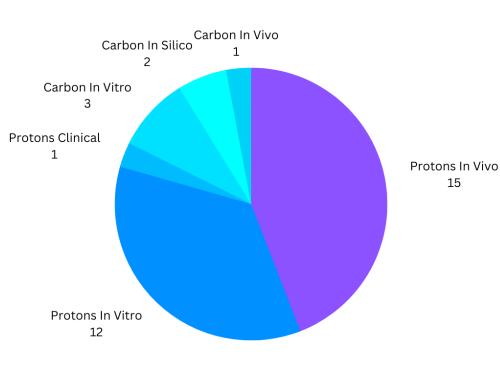
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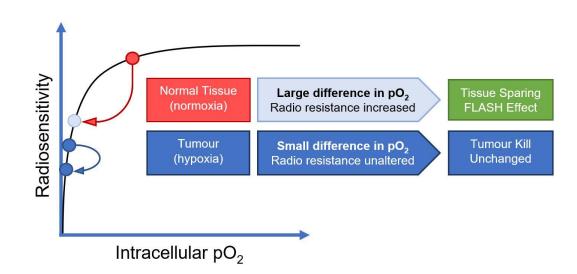
## **Key Findings**

- 1. In vivo proton therapy studies are most prevalent with a total count of 15
- 2. Six studies utilise the Bragg peak for irradiations, rest use transmission
- 3. Stringent verification of dosimetry is rarely provided, not yet standardised
- 4. Biological assays are prone to limitations which need greater acknowledgement
  - Atkinson, J., Bezak, E. & Kempson, I. Imaging DNA double-strand breaks — are we there yet?. Nat Rev Mol Cell Biol 2022
- 5. Mechanistic understanding of biological mechanisms is still in its infancy

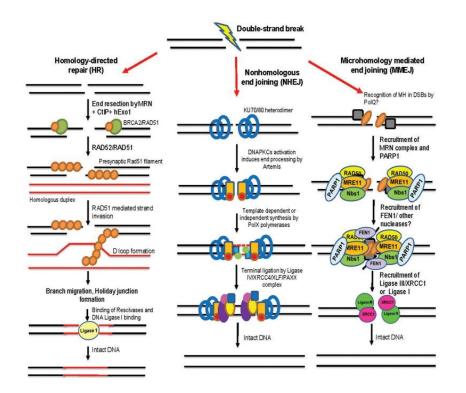
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### **Mechanism 1: Oxygen Depletion Hypothesis**

- Reactive oxygen species, produced by ionising radiation, cause indirect DNA damage
- Deplete O2 → Fewer ROS →
  Reduced damage
- Normal tissues large O2 change
- Tumours small O2 change







### Mechanism 2: DNA Damage and Repair

- <u>Differential effect</u> between normal and cancerous tissues may reside in how cells respond to DNA damage
- Hypothesised that FLASH causes DNA lesions of higher complexity than CONV
- Dysfunction of these pathways is common in cancers - inability for cancers to repair complex DNA lesions responsible?
- Capacity of normal cells to repair complex lesions better than cancerous cells?

Pandey and Raghavan 2017, J Radiat Cancer Res

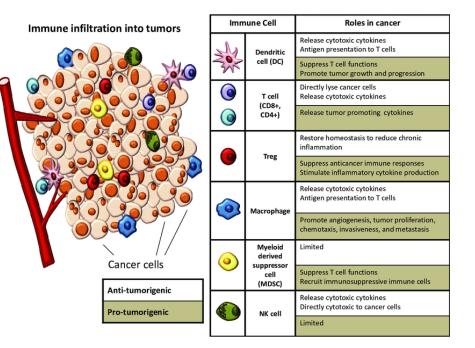
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### Mechanism 3: Immune System Modulation

- Anti-Cancer Effect Tumour Tissues
  - Improvement to tumour response and abscopal effect?
  - Improved immune cell influx, reduced vascular collapse in Lewis Cell Carcinoma in mice, Young-Eun Kim et al. (2021)
  - Dendritic, NK, T-Cells cytotoxic!
- Anti-Inflammatory Effect Normal Tissues
  - Does FLASH spare specific normal cell subpopulations (e.g. stem cells, immune cells)?
  - Proton-FLASH reduces TGFB1 expression (inflammatory marker) in normal cells Buonanno et al. (2019)
  - Simulations predict a reduction in circulating immune cells killed during FLASH (5-10%) vs low dose rate irradiation (90-100%), Jin, J. et al. (2020)



Markman and Shiao 2015, J Gastrointest Oncol



# Imaging DNA double-strand breaks — are we there yet?

Jake Atkinson<sup>1</sup>, Eva Bezak<sup>2,3</sup> and Ivan Kempson<sup>1</sup>

DNA double-strand breaks (DSBs) are fundamental to cell biology, from evolution to the latest gene-editing technologies. Yet, does an assay exist that truly quantitatively visualizes DSBs? Over-reliance on DSB detection by proxies can misguide interpretation of conventional assays, and more faithful DSB representatives await development.

DNA double-strand breaks (DSBs) can arise spontaneously from nuclease activities or through insults from reactive species, radiation and cytotoxic agents. DSBs are the most deleterious form of DNA damage, causing genome instability and chromosomal aberrations; DSB repair responses are thus rapid and stringently regulated<sup>1</sup>. By contrast, prescribed cleavage of DNA at targeted loci facilitates gene editing technologies such as CRISPR-Cas.

Quantifying DSBs is essential for identifying relationships between DNA cleavage and repair, which are pertinent to ageing, mutagenesis, genotoxicity and therapeutics, both established (chemo- and radio-therapies) and emerging (gene editing, particle and FLASH radiotherapy, and so on). Currently, our most powerful tools to spatially observe DSBs rely on immunofluorescence labelling of DNA damage response (DDR) proteins or histone modifications such as 53BP1 and γH2AX, respectively. These well-established markers typify the DDR<sup>2</sup>. However, interpretation of these downstream proxies is often extended upstream, through a range of assumptions, to DSB formation in a directly proportional, quantitative manner. These interpretations can be problematic: repair can be partial, thereby skewing

protein is 53BP1, which recruits additional DSB repair proteins and regulates repair-pathway choice. Through immunofluorescence, 53BP1 and  $\gamma$ H2AX are widely used, often dogmatically, as DSB markers, as they form discrete foci at or in proximity to DSBs.

#### **Biases of current assays**

The only conventional method for in situ DSB detection is terminal deoxynucleotidyl transferase dUTP nick-end labelling (TUNEL), but it does not quantify discrete DSB foci. Consequently, other markers are used, but considering many reports in the literature, it is necessary to remember that DDR markers generally reflect repair, not DSBs. An excellent example is perturbation of the DDR with a TGFB inhibitor prior to DNA insult. Consequently, fewer yH2AX foci are detected compared with control, which correlates with decreasing clonogenicity, that is, repair is impaired, leading to reduced proliferative capacity<sup>5</sup>. This highlights the crux of assays based on DDR factors: they are ambiguous. Do foci correlate with DSB accumulation, increased repair capacity and/or efficiency, or expression of DDR factors? Other misconceptions when using DDR proteins as DSB markers may arise.

## **Nature Comment**

- 'Mini review' accepted into Nature Reviews Molecular Cell Biology: Impact Factor = 113.9
- Imaging DSBs is useful for understanding the impact of cytotoxic agents i.e. radiotherapy
  - Primary mode of tumour cell death
- Current detection techniques lack target specificity they are proxies: proteins that signal *breakage repair*, but not the breaks themselves (γH2AX)
- Paper delves into why current assays are biased, novel alternatives, and advantages/disadvantages
- Desire for DSB-specific techniques
  - dSTRIDE currently optimising this technique to compare its effectiveness versus "conventional" DSB detectors



## **Future Experiments – In Vitro**

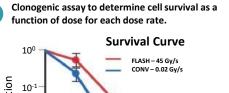
In vitro cell line irradiations with a range of doses, FLASH vs CONV. FLASH CONV 5 0 5 А 15 10 15 10 в 20 25 20 25 С

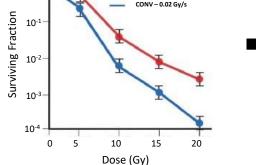
### Tumorigenic cell lines:

- 4T1 Breast Tissue
- K7, K7M2 Osteosarcoma

#### Normal cell lines:

- BALB/c Primary Lung Fibroblasts
- BALB/c Mouse Primary Bone Marrow-Derived Endothelial Cells





To determine extent of breaks 'clustering', foci will be quantified in MATLAB.

γH2AX/53BP1 + dSTRIDE immunofluorescence to compare DSB detection techniques + "quantify" DSBs...

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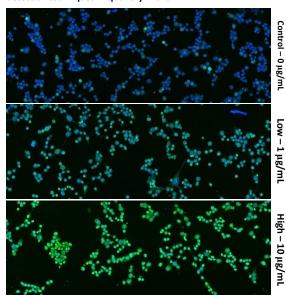


Figure: 4T1 cells vs varying concentrations of neocarzinostatin



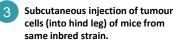
## **Future Experiments – In Vivo**

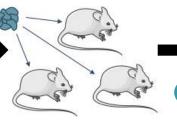
*In vivo* tumour from inbred strain BALB/c. Syngeneic model, <u>immunocompetent.</u>





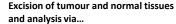
In vitro growth of tumour cells (4T1 and K7M2).





Irradiation of mice with either FLASH (45Gy/s) or CONV (<0.02Gy/s) protons. Tumour and comparative normal tissue irradiations will be performed.

Normal skin irradiations will also be scored qualitatively to record dermatitis severity i.e. erythema, desquamation, necrosis. 5 Exci and





- 1. Hematoxylin and Eosin (H&E), Masson Trichrome, TUNEL Staining
  - Tissue structure and morphology
  - Immune cell infiltration
  - Apoptotic cells

#### 2. qPCR, Western Blot

- Aim to quantify markers of inflammation including:
- TGF- β
- IL-1, IL-6
- TNF-α
- NF-кВ
- MAPK, JAK-STAT pathways

#### 3. Flow Cytometry

- Immune cell population
- Tumour and normal cell
  microenvironment modulation





# Acknowledgements

- UniSA, Future Industries Institute
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- MITRU, SAHMRI
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- PhD Supervisors Ivan Kempson, Eva Bezak and Hien Le
- Members of Kempson and Bezak Research Groups

### Thank you for listening!





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