

Benchmarking simulations by measured yields of initial DNA damage and late effects for ion beam irradiation of human and mammal cells.

Faire avancer la sûreté nucléaire

IRSN/PRP-HOM/SRBE/LDB (France)

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ENEA (Italy)

C. PATRONO, A. TESTA, M. PINTO

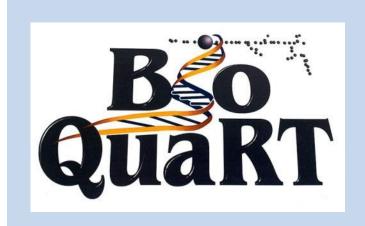
IST/ITN (Portugal)

O. MONTEIRO GIL

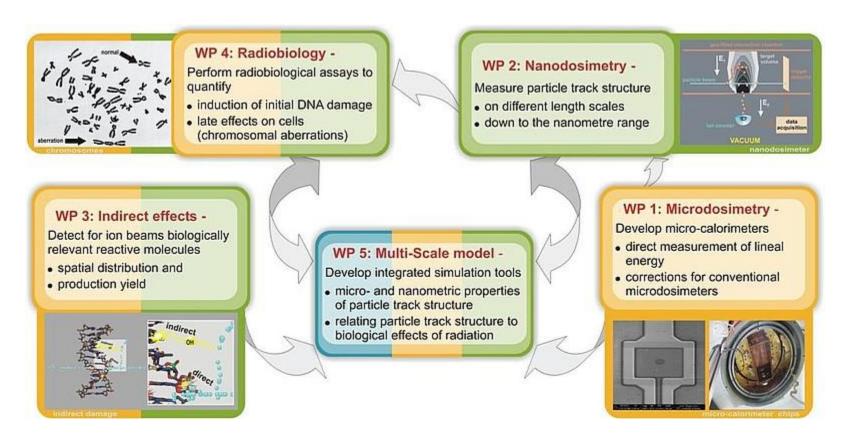
PTB (Germany)

U. GIESEN

BioQuaRT Midterm Dissemination Workshop 7 June 2013

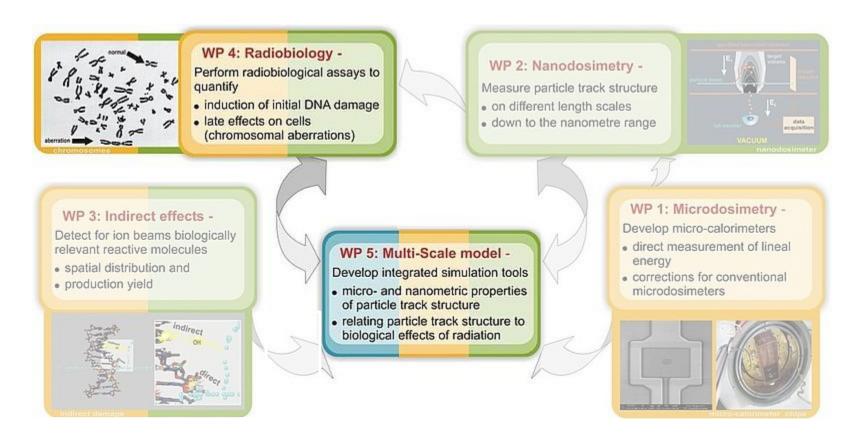


WP4: Radiobiology



Work package 4 focuses on the biological aspects of radiation damage and will produce data on initial DNA damage and late effects for ion beams of different radiation qualities.

WP4: Radiobiology



The biological data collected in this WP will be used for benchmarking with the multi-scale model developed in WP5.

Challenges of WP4

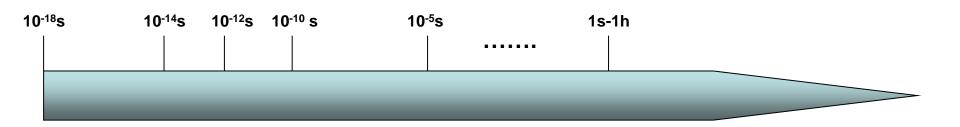
BioQuaRT Midterm Dissemination Workshop - 7 June 2013

■ To perform biological measurements that could "feed" the simulation

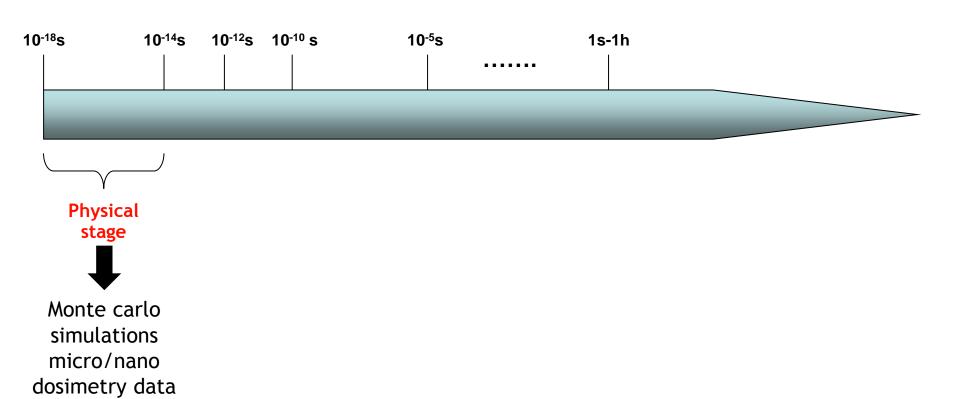
Challenges of WP4

- To perform biological measurements that could "feed" simulations
 - 1. To choose relevant biological endpoints

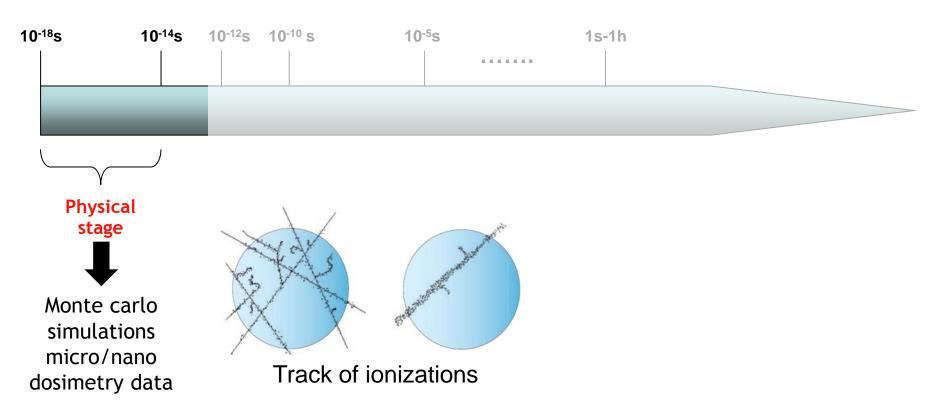
- Link between physics events and biological events
 - Post-irradiation timeline



- Link between physics events and biological events
 - Post-irradiation timeline

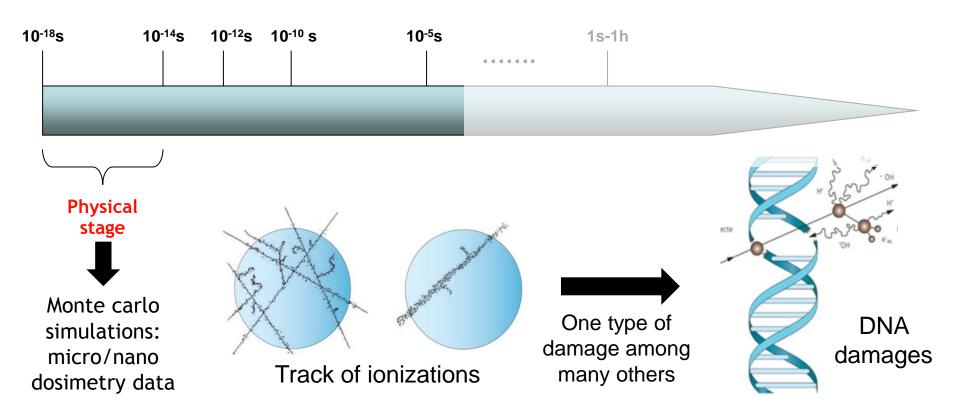


- Link between physics events and biological events
 - Post-irradiation timeline

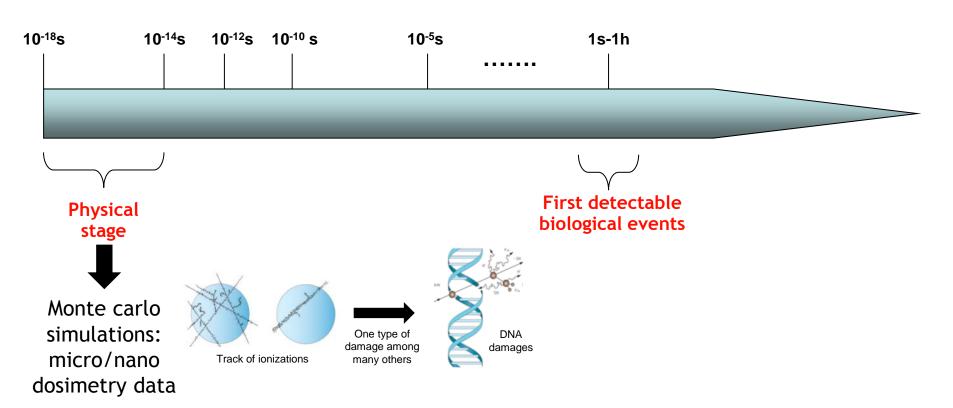


Link between physics events and biological events

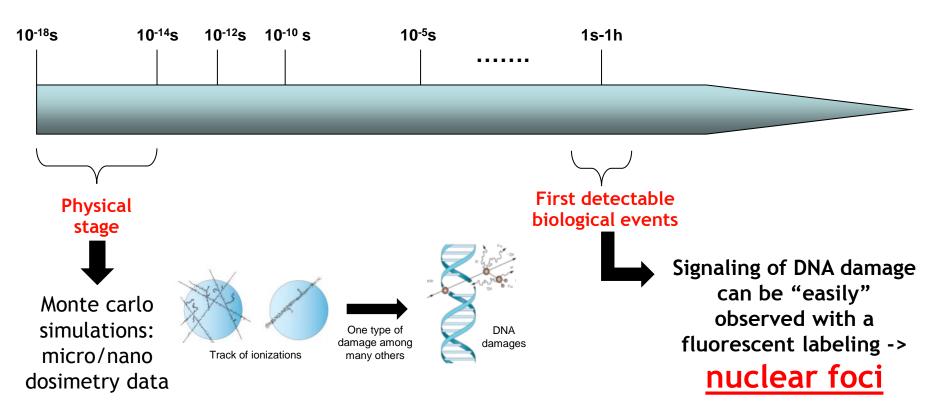
Post-irradiation timeline



- Link between physics events and biological events
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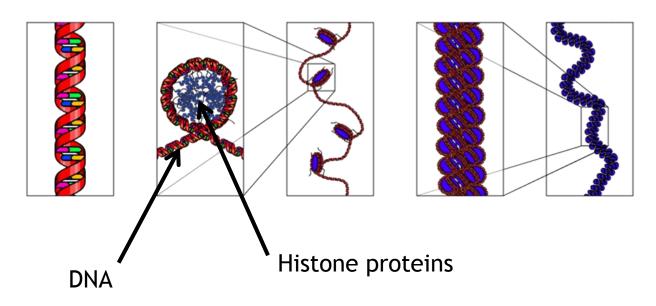
- Link between physics events and biological events
 - Post-irradiation timeline



Performed at IRSN (G. Gonon, Pa. Voisin and G. Gruel)

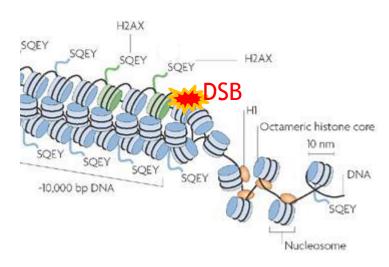
Nuclear foci

In the cell nucleus, the DNA molecule is compacted around proteins named histones



- Performed at IRSN (G. Gonon, Pa. Voisin and G. Gruel)
 - Nuclear foci

When a double strand break occurs in the DNA molecule

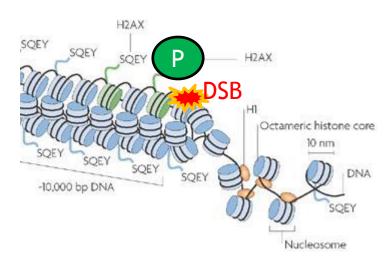


Bonner et al 2008

Performed at IRSN (G. Gonon, Pa. Voisin and G. Gruel)

Nuclear foci

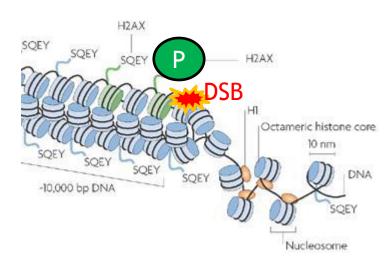
Histone proteins can be modified around the damage (such as γ-H2AX foci) or certain proteins are relocalized at the site of DNA DSBs (such as 53BP1)



Bonner et al 2008

- Performed at IRSN (G. Gonon, Pa. Voisin and G. Gruel)
 - Nuclear foci

And this modification and/or relocalization can be detected through a fluorescent labelling

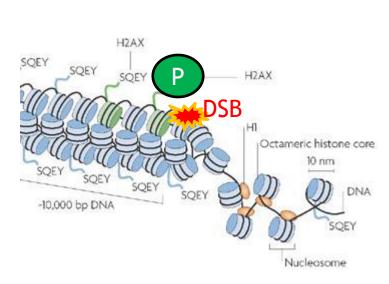


Bonner et al 2008

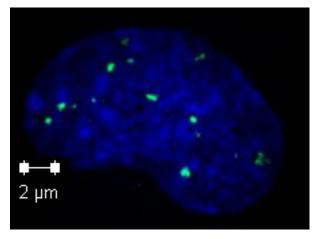
Performed at IRSN (G. Gonon, Pa. Voisin and G. Gruel)

Nuclear foci

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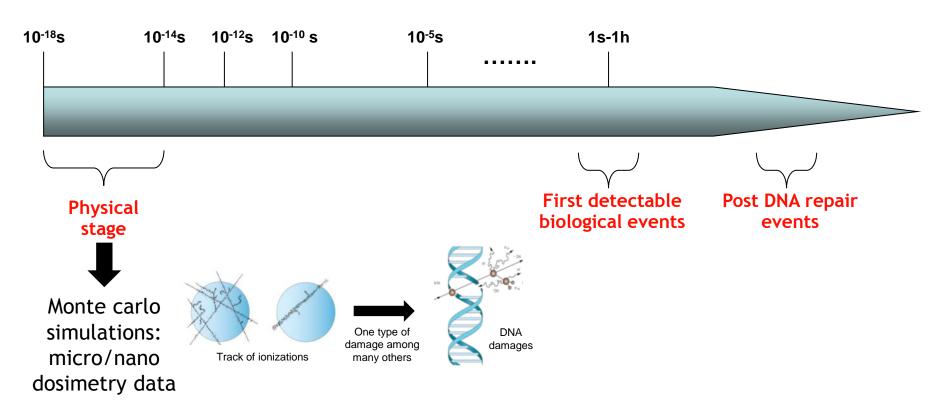


Bonner et al 2008

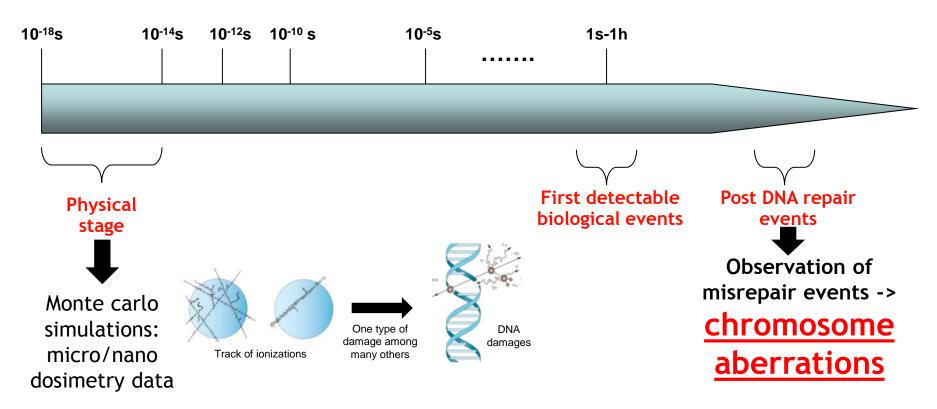


We can then analysed quantity, quality and topology of these foci

- Link between physics events and biological events
 - Post-irradiation timeline



- Link between physics events and biological events
 - Post-irradiation timeline



Biological endpoint: late events

- Performed at ENEA (C. Patrono, A. Testa) and IST/ITN (O. Monteiro Gil)
 - Chromosome aberrations

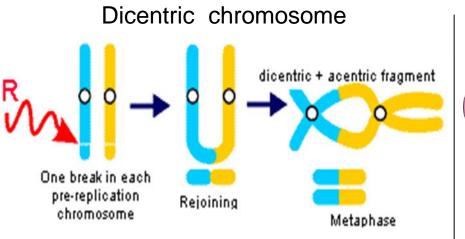
The DNA ends may rejoin in different patterns from their original arrangement. The abnormalities that result are termed "chromosome aberrations" (CA) and may be visualized in specific stages of mitosis.

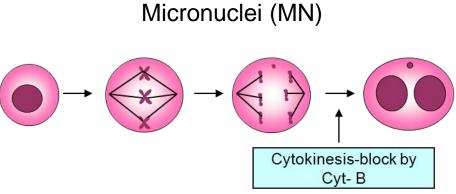
The frequency of CA is known to change with radiation dose and/or quality.

Biological endpoint: late events

- Performed at ENEA (C. Patrono, A. Testa) and IST/ITN (O. Monteiro Gil)
 - Chromosome aberrations

Two kinds of CA will be scored:

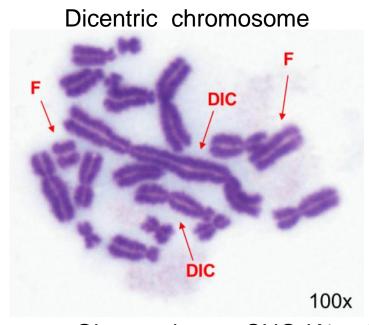


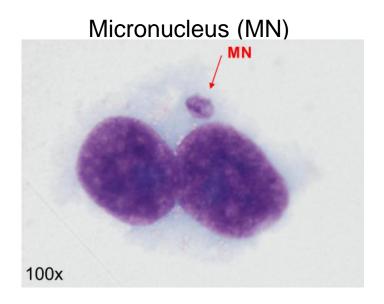


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Two kind of CA will be scored:





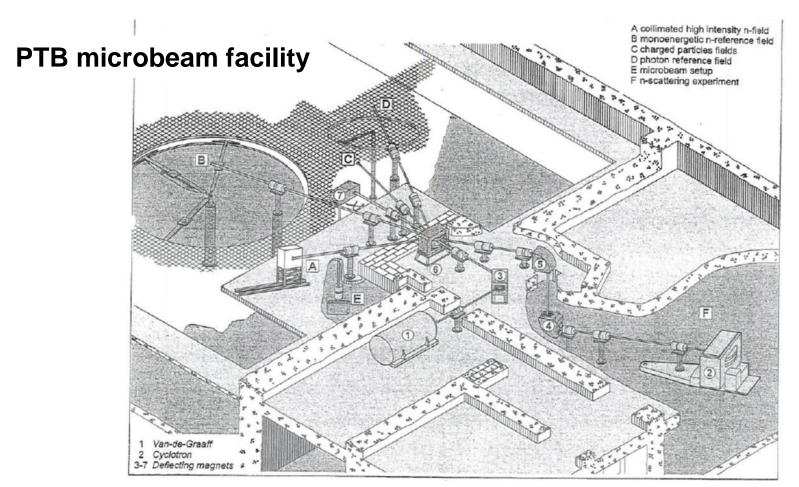
Observation on CHO-K1 cells after exposure to 20 MeV α particles

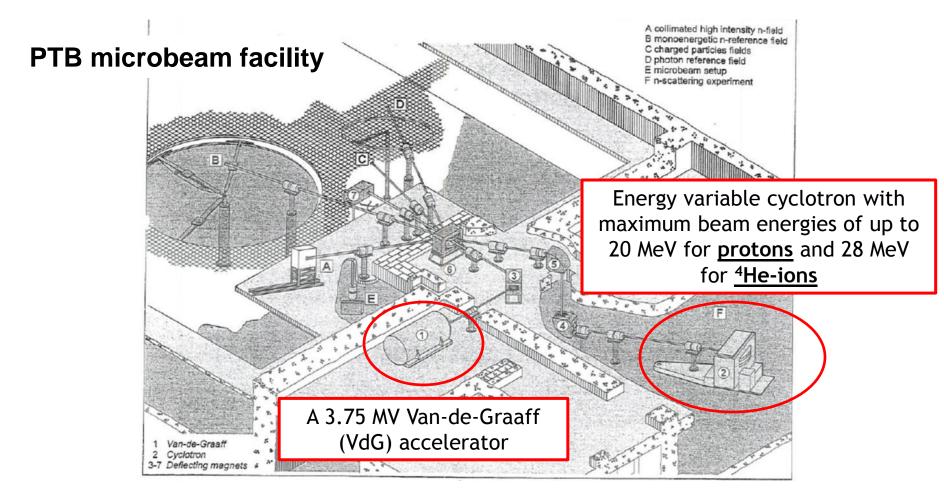
Challenges of WP4

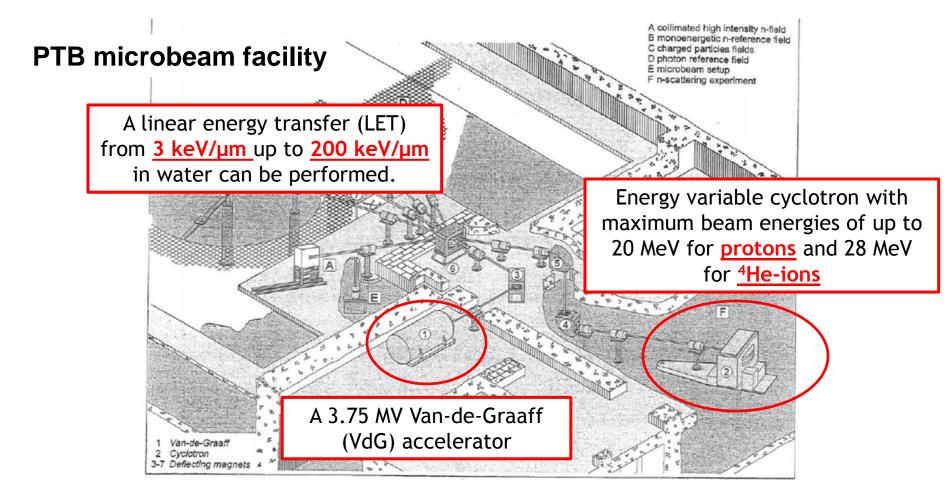
- To perform biological measurements that could "feed" WP5
 - 1. To choose relevant biological endpoints

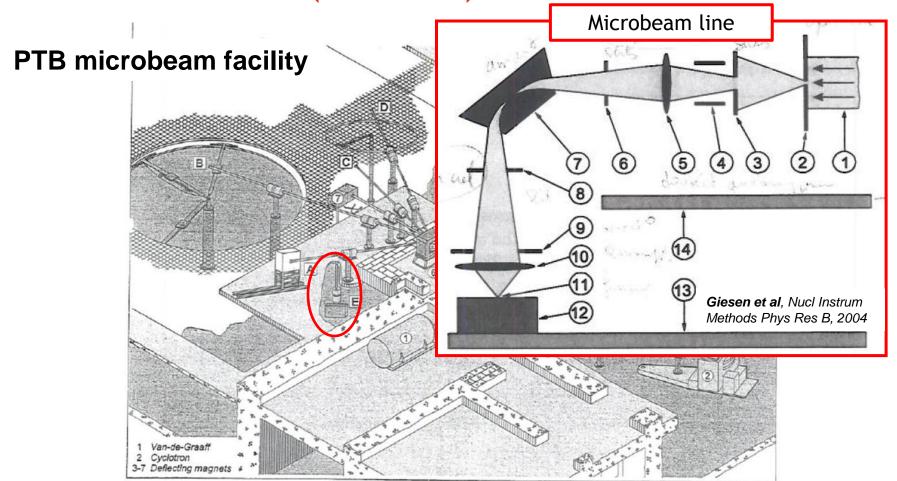
Challenges of WP4

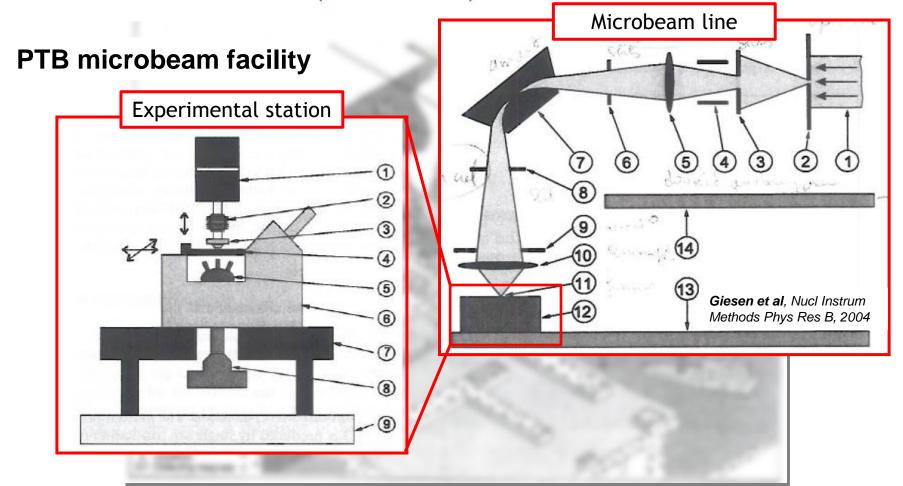
- To perform biological measurements that could "feed" WP5
 - 1. To choose relevant biological endpoints
 - 2. To choose relevant irradiation conditions...

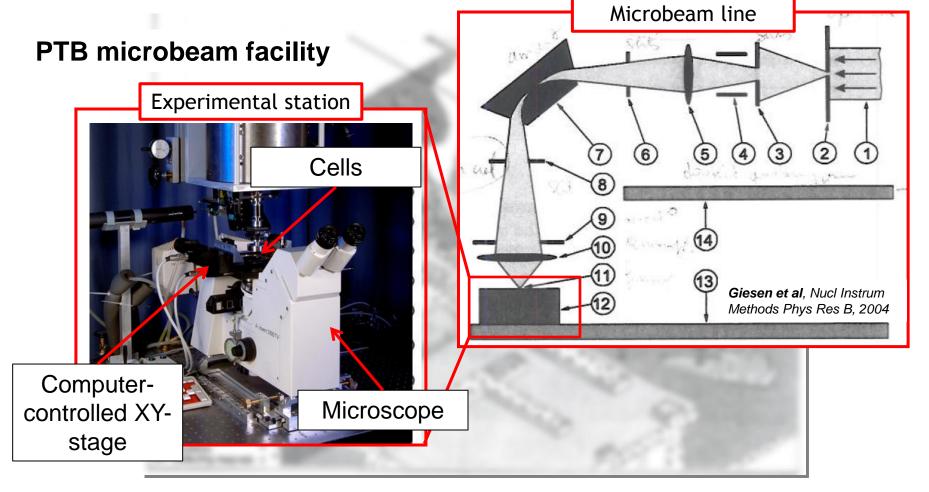






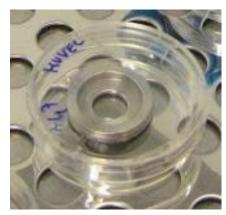






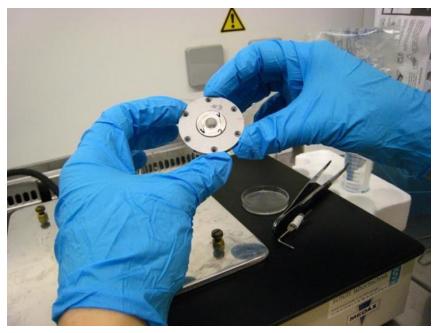
Cell are plated in special dishes the day before





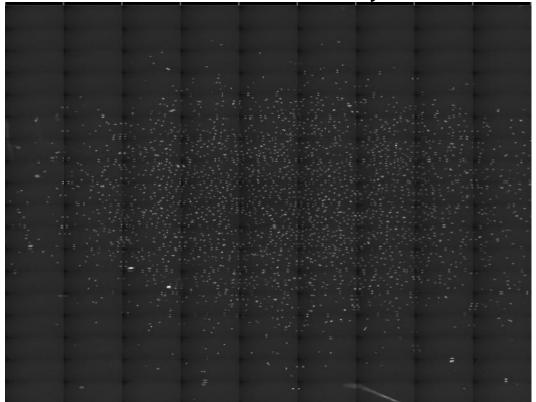
- The day of irradiation, cells are stained with Hoechst (Nucleus staining)
- Then, the dish is mounted in a special device

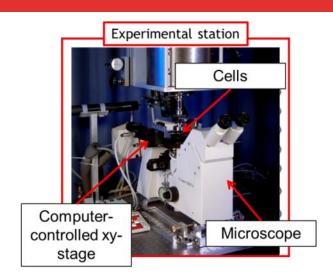




Detection of cell nuclei

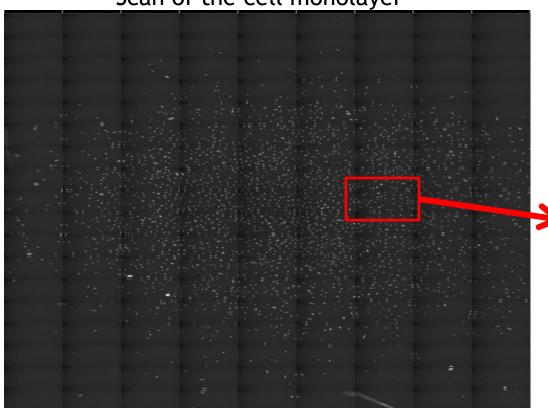
Scan of the cell monolayer

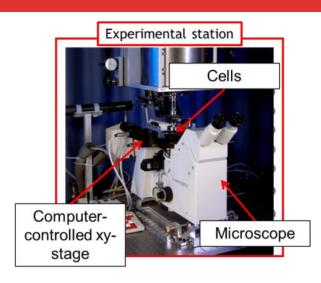


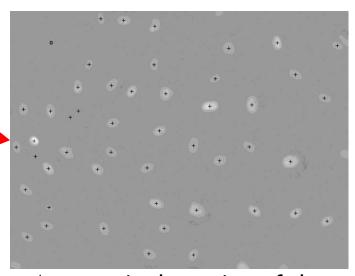


Detection of cell nuclei

Scan of the cell monolayer

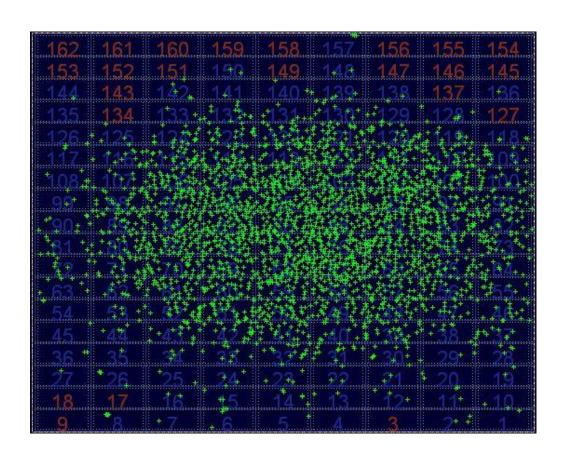


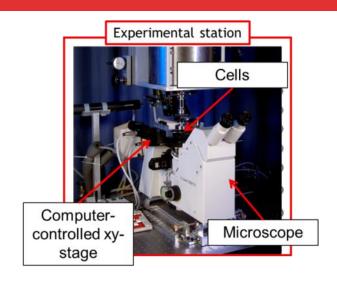




Automatic detection of the barycenter of each nucleus of the cell population

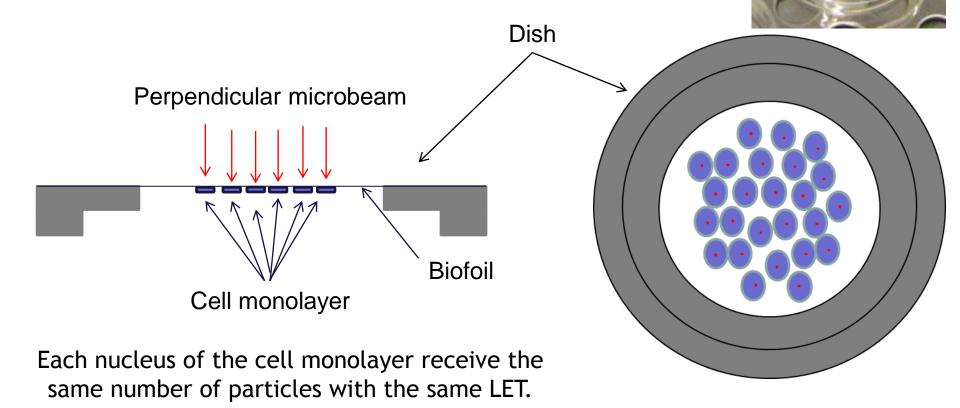
Detection of cell nuclei





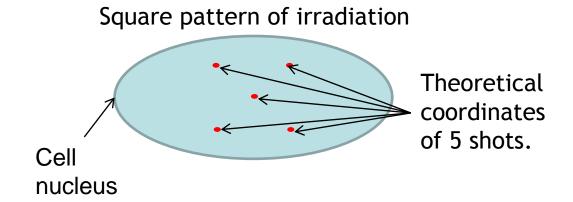
The coordinates of the "microbeam irradiation" are then computed.

Geometry of the microbeam irradiations



Types of irradiation already performed for "early events"

Alpha	Proton
8 MeV (160 keV/μm)	3 MeV (18 keV/μm)
20 MeV (37 keV/μm)	10 MeV (5 keV/μm)



Challenges of WP4

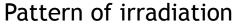
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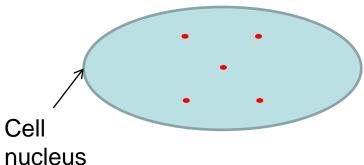
Challenges of WP4

- To perform biological measurements that could "feed" WP5
 - 1. To choose relevant biological endpoints
 - 2. To choose relevant irradiation conditions...
 - 3. ...to be able to perform relevant measurements.

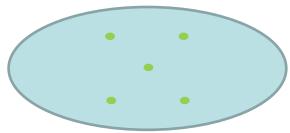
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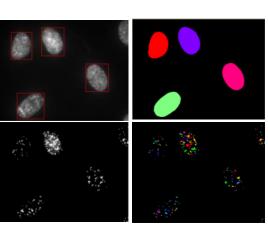
How many of resulting foci by cell nucleus?



Performed at IRSN (G. Gonon, Pa. Voisin and G. Gruel)

Mass analysis





With the combination of

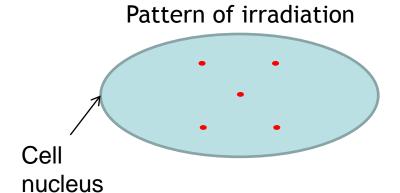
- high speed microscopy platform
- Automated image analysis

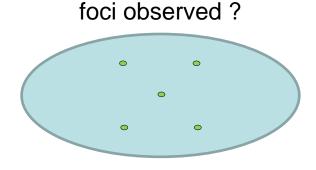
We can analyse quantity, quality and topology of foci on a large population of nuclei (>2000) exposed to the same irradiation condition

This provides a measure of the probability of the presence (or absence) of foci within a cell nucleus.

Types of irradiation already performed for "early events"

Alpha	Proton
8 MeV (160 keV/μm)	3 MeV (18 keV/μm)
20 MeV (37 keV/µm)	10 MeV (5 keV/μm)

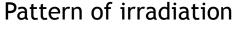


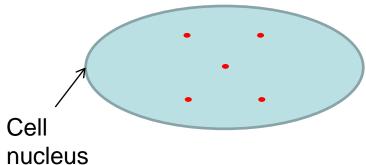


for ex: probability of hit: 5/5

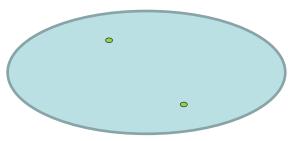
Types of irradiation already performed for "early events"

Alpha	Proton
8 MeV (160 keV/μm)	3 MeV (18 keV/μm)
20 MeV (37 keV/μm)	10 MeV (5 keV/μm)





foci observed?



for ex: probability of hit: 2/5

DAPI 53BP1 (Tx-Red) γ-H2AX (FITC) Merged

5α-irradiated dish

(20 MeV, 37keV/µm)

DAPI 53BP1 (Tx-Red) γ-H2AX (FITC) Merged

5α-irradiated dish

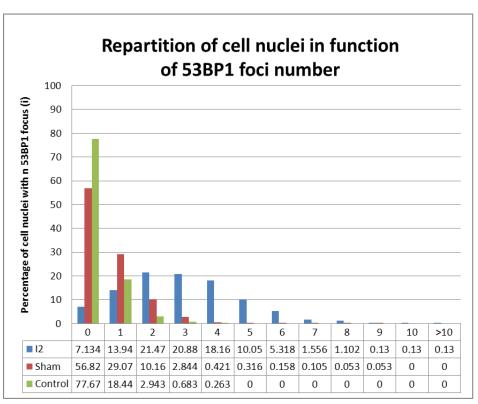
(20 MeV, 37keV/µm)

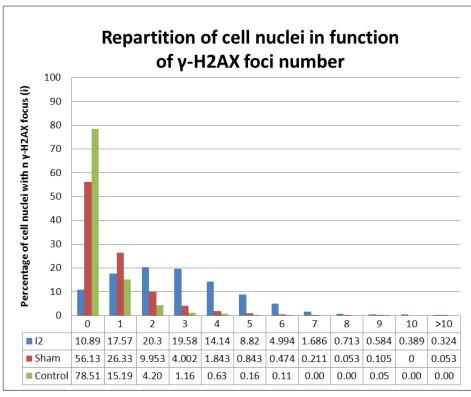
DAPI 53BP1 (Tx-Red) γ-H2AX (FITC) Merged

5α-irradiated dish (20 MeV, 37keV/μm)

Mass Analysis

7 HUVEC 5 alpha particles pattern (20 MeV, 37 keV/μm), Nov 28th, 2012 - more than 3000 analyzed





Challenges of WP4

- To perform biological measurements that could "feed" WP5
 - 1. To choose relevant biological endpoints
 - 2. To choose relevant irradiation conditions...
 - 3. ...to be able to perform relevant measurements.

Challenges of WP4

- To perform biological measurements that could "feed" WP5
 - 1. To choose relevant biological endpoints
 - 2. To choose relevant irradiation conditions...
 - 3. ...to be able to perform relevant measurements.
 - 4. To take into account the "uncertainties" of irradiation conditions and biological models

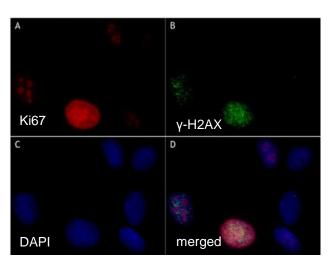
- Biological parameters
 - Foci background

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 - the stage of the cell cycle

- Biological parameters
 - Foci background
 - the stage of the cell cycle

Cells in phase S, G_2 or M can be excluded from the analysis using a combination of several parameters measured on each nucleus as:

- Integrated fluorescence of DAPI (related to DNA quantity) and
- Integrated fluorecence of FITC (related to phosphorylation of H2AX due to DNA synthesis)





- Biological parameters
 - Foci background
 - the stage of the cell cycle SOLVED -
 - Interaction between Hoechst staining (nuclei stain) and UVscan

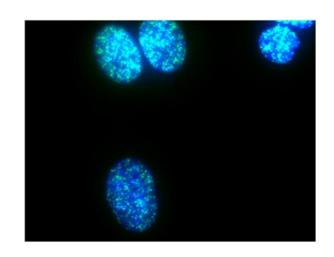
Biological parameters

- Foci background
 - the stage of the cell cycle SOLVED -
 - Interaction between Hoechst staining (nuclei stain) and UVscan

Illumination of live cells labelled with Hoechst produces y-H2Ax foci

This was observed when Hoechst was excitated with a **mercury lamp** combined with adequate filter.

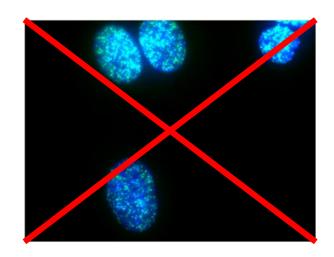
This effect depends on **time** of illumination and **concentration** of Hoechst



Biological parameters

- Foci background
 - the stage of the cell cycle SOLVED -
 - Interaction between Hoechst staining (nuclei stain) and UVscan

The replacement of the mercury lamp with new LED light source (Lumencor Spectra X) seems to totally remove the effect.



- Biological parameters
 - Foci background
 - the stage of the cell cycle SOLVED -
 - Interaction between Hoechst staining (nuclei stain) and UVscan - SOLVED-

Biological parameters

Foci background

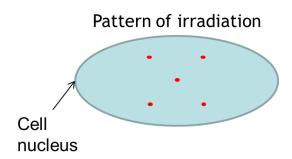
- the stage of the cell cycle SOLVED -
- Interaction between Hoechst staining (nuclei stain) and UVscan - SOLVED-

Beam parameters

- Beam size
- Scintillator noise



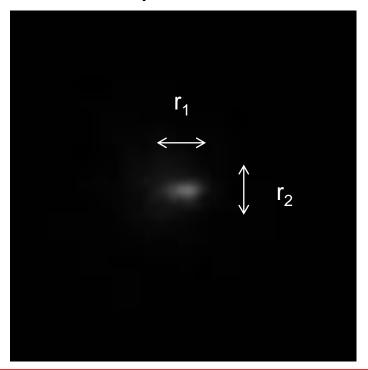
- Biological parameters
 - Foci background
 - the stage of the cell cycle SOLVED -
 - Interaction between Hoechst staining (nuclei stain) and UVscan - SOLVED-
- Beam parameters
 - Beam size
 - Scintillator noise



Could lead to the deformation of the theorical pattern of irradiation

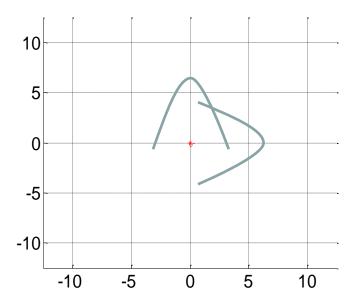
Beam parameters

- Beam size: straddling effect and beam focalization
 - This could be estimated => the beam shape is measured for each microbeam set-up.



Beam parameters

- Beam size: straddling of different layers and beam focalization
 - Coordinates of each hit of the pattern is a function of a Gaussian distribution in X and Y (μ =0, σ =r1/2.32 and σ =r2/2.32)



Beam parameters

Scintillator Noise

- The noise of scintillator could be interpreted as a particle. This leads to the closure of the beam shutter even if no particle has been emitted. As a consequence no particle can be emitted instead of 1.
- At the contrary, the path of one particle through the scintillator could be interpreted as a noise. This leads to the non-closure of the beam shutter leaving another particle coming. As a consequence 2 particles can be emitted instead of 1.

Signal of the scintillator is saved for each irradiation: the overlap between background noise and real signal is estimated.

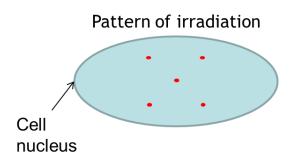
Beam parameters

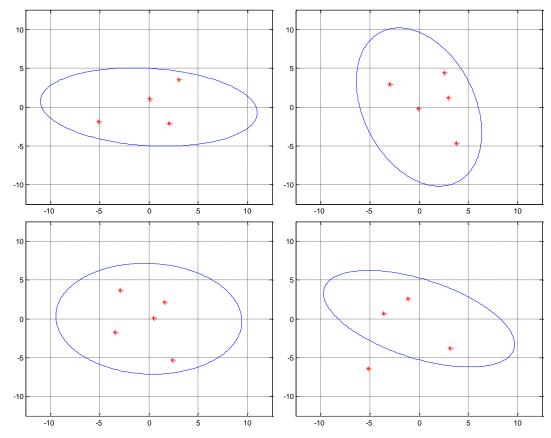
- We develop a script with Matlab to simulate how
 - The beam size
 - The scintillator noise
 - The nucleus dimension

can affect the initial pattern of irradiation and then the foci distributions observed



Beam parameters





Biological parameters

Foci background

- the stage of the cell cycle SOLVED -
- Interaction between Hoechst staining (nuclei stain) and UVscan - SOLVED-

Beam parameters

- Beam size
- Scintillator noise

- Effects can be estimated -

Conclusion

It seems possible to remove at least a portion of the noise of the observed foci distribution among cell population.

The objective is to access to an accurate estimation of the probability of interaction between a given particle with a given LET and DNA.

With these kind of corrected measurements, it will be possible to compare biological results with Monte Carlo simulation by benchmarking.

Thank you