

# DEVELOPMENT OF A TIME-OF-FLIGHT COMPTON CAMERA FOR ONLINE CONTROL OF ION THERAPY

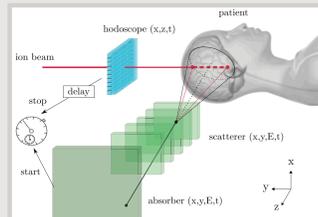
J.-L. Ley<sup>1</sup>, C. Abellan<sup>2</sup>, J.-P. Cachemiche<sup>2</sup>, M. Dahoumane<sup>1</sup>, D. Dauvergne<sup>1</sup>, N. Freud<sup>3</sup>, B. Joly<sup>4</sup>, J. Krimmer<sup>1</sup>, L. Lestand<sup>4</sup>, J.M. Létang<sup>3</sup>, H. Mathez<sup>1</sup>, V. Maxim<sup>3</sup>, G. Montarou<sup>4</sup>, C. Ray<sup>1</sup>, M-H Richard<sup>1</sup>, E. Testa<sup>1</sup>, Y. Zoccarato<sup>1</sup>

<sup>1</sup> Institut de Physique Nucléaire de Lyon; <sup>2</sup> CPPM, Aix-Marseille Université, CNRS/IN2P3, Marseille; <sup>3</sup> Laboratoire CREATIS, Lyon; <sup>4</sup> Laboratoire de Physique Corpusculaire de Clermont-Ferrand

## COMPTON CAMERA

### PURPOSE

The aim of irradiation monitoring during a treatment in ion therapy is to control in real time the agreement between the delivered dose and the planned treatment. In fact, the discrepancies might come from uncertainties such as the positioning or the morphological changes of the patient. They can lead to ion-range variations of a few millimeters. Several devices are under development over the world to detect secondary radiations, which are correlated to the dose deposited by incident ions. Compton cameras are in particular investigated for their potential high efficiency to detect prompt gamma rays [1,2,3,4]. The present work aims at discussing the clinical applicability of a Compton camera design by means of Monte Carlo simulations for proton and carbon ion therapy.



**Fig. 1.** Configuration of the monitoring system: the prompt gamma-ray emission points are reconstructed by intersecting the ion trajectory and the Compton cone. The ion trajectory is obtained with the hodoscope and the Compton cone is reconstructed with the camera. Time of flight (TOF) measurements between the absorber detector and the beam hodoscope (with an appropriate delay) are performed. [5]

## SIMULATIONS GEANT4.9.6

### KEY POINTS

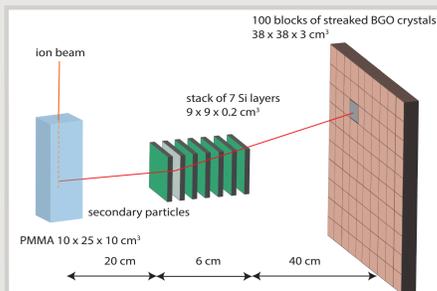
- Monte-Carlo simulation: Geant4.9.6
- Use the standard electromagnetic processes and the Binary Cascade model for hadrons
- Taking account of the Doppler broadening and polarization
- Cut at 0.1 mm in PMMA and 1 mm in the detectors

	Protons	Carbon ions
Number	$10^8$	$4 \times 10^7$
Energy	160 MeV	305 MeV/u
Bragg peak position	15.5 cm	

**Table 1.** Parameters used for the simulation of the Compton camera.

Resolution (FWHM) @ 1 MeV	BGO	Si
Spatial [mm]	5	0.9
Energy [%]	17	2.3
Timing [ns]	3	15

**Table 2.** Realistic resolutions defined in the simulations for the detectors. The energy resolution for the silicon is defined with the Equivalent Noise Charge (ENC). The ENC for the silicon is expected to be 200.



**Fig. 2.** Diagram of the simulation setup.

## DATA TREATMENT

A beam time structure is modeled and applied to the simulation data in order to estimate the coincidence rate (true/random) for a clinical application of the Compton camera. The treatment is done for protons and carbon ions.

### CLINICAL CHARACTERISTICS

	Protons	Carbon ions
Facility	Cyclotron C230 IBA	Synchrotron at HIT
Clinical intensity	$2 \times 10^{10}$ p/s	$5 \times 10^7$ ions/s
Energy	160 MeV	305 MeV/u
Bunch time [ns]	3.2	30
Period [ns]	9.4	170
Particles / bunch	217	9
Coincidence window [ns]	40	40

### BEAM STRUCTURE

### DETECTORS

**Table 3.** Characteristics of clinical beams using protons and carbon ions. Each particle is treated as an individual story in the simulation Geant4. The beam time structure is modeled in the simulation data analysis.

## RESULTS

	Structure	COINCIDENCE RATE FOR PROTONS						COINCIDENCE RATE FOR CARBON IONS				
		Single proton		IBA				Single ion		HIT		
		Intensity	10 <sup>6</sup> p/s	1.2 x 10 <sup>9</sup> p/s	2 x 10 <sup>10</sup> p/s	perfect	real	perfect	real	perfect	real	
Coincidence [%]	Detectors	perfect	real	perfect	real	perfect	real	perfect	real	perfect	real	
	Random	0	0	0.3	47.1	7.1	92.3	≈ 0	$8.9 \times 10^{-2}$	4.7	64.0	
	all	100	100	99.7	52.9	92.9	7.7	≈ 100	99.9	95.3	36.0	
	True	same gamma	92.0	51.5	91.7	27.3	85.5	2.9	84.6	22.5	80.6	7.7
		other	8.0	48.5	8.0	25.6	7.4	3.8	15.4	77.4	14.7	28.3
Count rate coincidence [MHz]		/	/	$7.7 \times 10^{-2}$	0.2	4.7	15.3	/	/	0.03	0.3	
Count rate single [MHz]	1 <sup>st</sup> silicon layer	/	/	1.4	22.7	/	/	/	/	1		
	absorber	/	/	16.8	281	/	/	/	/	8.1		

**Table 4.** The relative coincidence rates are given for three different proton and carbon ion intensities. There are no cuts in energy or with time of flight. The count rate is the sum of the random and the true coincidences. The count rate single is the number of particles which deposit energy in the detector independently of a coincidence. The first silicon layer has the highest single count rate.

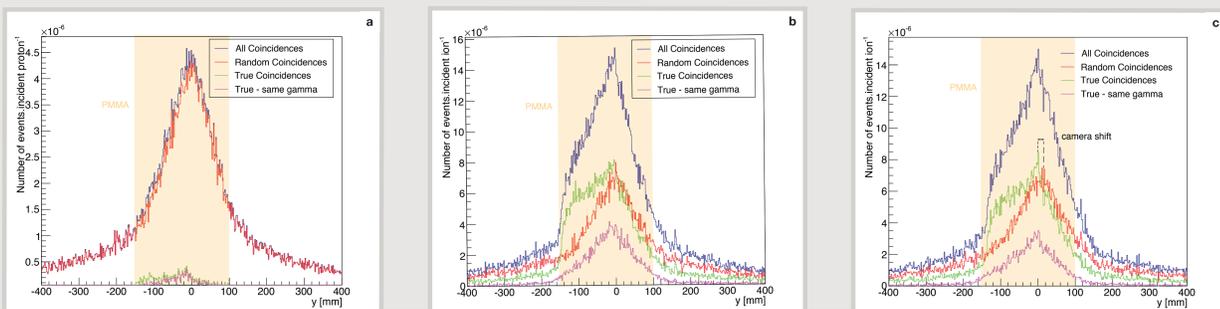
### DEFINITIONS

#### STRUCTURE

- **Single particle:** One particle is run each microsecond
- **IBA or HIT:** Application of the specific time beam structure

#### DETECTORS

- **Perfect:** no timing resolution is applied. The time window defined to detect a coincidence is the time for a photon to fly from the silicon layer to the absorber.
- **Real:** a realistic timing resolution is applied to the detectors (table 3) and the time window for a coincidence is set at 40 ns.



**Fig. 4.** Reconstructed profile in the case of protons (a) and carbon ions (b,c) with the time structure of the beam and realistic detector resolutions. The theoretical falloff position is 0 mm along the Y axis. For the figure 4a and 4b, the camera is centred on the Bragg peak position. In the figure 4c, the camera is shifted to 50 mm after the Bragg peak position. It shows that the maximum of the random coincidence distribution is due to the position of the camera and not related to the falloff position. Therefore, we have to distinguish true coincidences which are the only way to get the falloff position.

## DISCUSSION

### PROTONS

- The fraction of random coincidences is about 90 % with a clinical beam and realistic detectors. This can be explained by:
  - **The beam structure:** The number of protons per bunch at  $2 \times 10^{10}$  p/s is 217. Therefore, there is a large amount of secondary particles at the same time that contribute to the random coincidences.
  - **The timing resolution of silicon:** The 15 ns induce a large uncertainty on the real interaction time and require a large coincidence window.
- Possibilities to improve the rate of true coincidences:
  - **Cuts in deposited energy:** An analysis on the deposited energies in the detectors was made in order to eliminate the coincidences due to charged particles. It works well for high energy (deposited energy >10 MeV in BGO and > 1 MeV in Si). However, the ratio between true and random coincidences is not improved because the majority of the random coincidences deposit a small amount of energy. It appears that the energy selection is not sufficient to improve the fraction of true coincidences.
  - **Use of the time of flight information:** The time of flight between the hodoscope and the absorber (or using the cyclotron HF signal) could help to differentiate gamma which are faster than neutrons or protons.
  - **Smaller coincidence window:** The fraction of random coincidences reduced to ~10% with a perfect detectors which let some perspectives for Compton camera using very fast detectors.
  - **Increase distance:** at 1 meter, the random coincidences are divided by a factor 50, whereas the true coincidences are only divided by 10. However, with only  $1.1 \times 10^9$  true coincidences per second, we will no longer be able to reach the desired statistics for a beam spot.
- **Conclusion:** For a clinical beam intensity ( $2 \times 10^{10}$  p/s) with these detectors and for this setup, the number of protons in a bunch is too high. The true coincidences are completely dominated by the random coincidences. A solution could be, at least, to decrease the intensity for the distal spots and then to have a higher intensity beam for the rest of the tumor. With the diminution of the clinical intensity, the quality assurance could be possible in real time with this Compton camera.

### CARBON IONS

- The coincidence rate is more favorable for carbon ions thanks to the lower beam intensity.
- The data analysis is still under progress. The number of neutrons and protons generated by nuclear reactions is high. Therefore, we expect that the time of flight information will improve the rate of true coincidences.
- The Compton camera applicability for carbon ion therapy seems more realistic.

### RECONSTRUCTION

With a standard MLEM reconstruction algorithm, the source distribution is reconstructed by intersecting all the Compton cones together. Some of the cone intersections do not correspond to an actual source position. These points are eliminated by the iteration process. This should improve the reconstruction of the falloff position [6].

## PROTOTYPE DEVELOPMENT

### THERE ARE THREE PARTS:

#### 1. Hodoscope → Poster 32 (J. Krimmer)

- Scintillating square fibers:  $140 \times 1 \times 1$  mm<sup>3</sup>
- Number of fibers:  $2 \times 128$
- 8 PMTs Hamamatsu multianodes H8500
- Timing resolution: 1 ns
- Count rate:  $10^9$  counts/s

#### 2. Scatterer (Double-sided silicon strip detector)

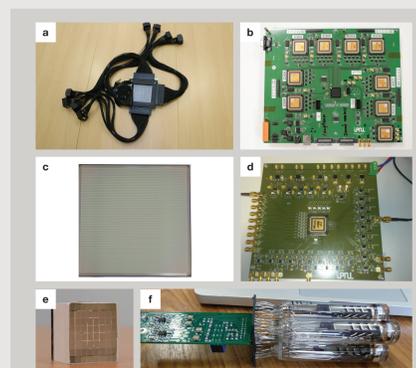
- Number of layers: 7
- Dimensions:  $90 \times 90 \times 2$  mm<sup>3</sup>
- Number of strips:  $2 \times 64$
- Pitch: 1.4 mm
- Expected performances
  - Spatial resolution: 1 mm
  - Energy resolution: 2 keV (FWHM) @ 1 MeV

#### 3. Absorber (streaked BGO)

- Dimensions:  $380 \times 355 \times 30$  mm<sup>3</sup>
- Number of blocks: 96
- 4 PMTs per block
- Timing resolution: 3 ns
- Spatial resolution: 4 mm

#### 4. μTCA acquisition system → Poster 3 (C. Abellan)

- Communication between the μTCA and the acquisition computer thanks to a gigabit ethernet link.



**Fig. 5.** a. The hodoscope with its 512 optical fibers connected b. Front End card for a first hodoscope prototype  $2 \times 32$  fibers c. Double-sided silicon strip detector d. First prototype of an ASIC for the DSSD's front end e. BGO block (96 blocks compose the absorber) f. Each block is connected to 4 PMTs

## REFERENCES

- [1] S.W. Peterson et al., Phys. Med. Biol. 55 (2010) 6841–6856
- [2] G. Llosa et al., NIMA 695(2012)105–108
- [3] T. Kormoll et al., IEEE-TNS (2011) 3484 - 3487
- [4] S. Kurosawa et al., Current Applied Physics 12 (2012) 364–368
- [5] M.-H. Richard et al., IEEE-NSS/MIC (2011) 3496–3500
- [6] X. Lojaco et al., IEEE-TNS (2013) 3355 - 3363

## CONTACT



Jean-Luc Ley  
PhD Student  
Mail: ley@ipnl.in2p3.fr  
Tel: +33 (0)4 72 43 14 81