# Monitoring ion beam therapy with a Compton Camera: simulation studies of the clinical feasibility

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Abstract. Technological progress and new experimental evidences guaranteed in the last years an important spread of particle therapy worldwide [Particle Therapy Cooperative Group, 2017]. Anyway, the present clinical routine is not yet completely exploiting the benefits of this treatment technique, due to treatment planning and delivery incertitude [Paganetti, 2012]. In order to further advance in this promising field, the need of a treatment monitoring method, ideally in realtime, results undeniable. Several techniques have been proposed, most of them based on the detection of secondary radiation produced by the beam interaction in the patient body [Knopf and Lomax, 2013, Parodi, 2016]. Among them, the detection of promptgamma rays has been deeply investigated in the last year [Krimmer et al., 2013], and many detectors are under construction by different research group in the world, in order to exploit spatial, energy and time information carried by this secondary particles. In particular, by detecting and analyzing the three variables at the same time, the beam depth-dose distribution can be retrieved by imaging techniques. Two main kind of detectors are suitable for this application: gamma cameras with mechanical collimation and Compton cameras. The ClaRyS French collaboration is developing a Compton camera prototype based on silicon detectors and BGO scintillator blocks [Krimmer et al., 2015], and the present work aims at discussing the clinical applicability of this design by means of Monte Carlo simulations. The Compton camera performance have been studied for different configurations: the Compton camera absolute efficiency varies in the range  $1-4 \times 10^{-4}$  with energy variations in the prompt gamma energy range; the number of detected coincidences as a function of the beam intensity shows that a beam intensity reduction is needed with respect to standard clinical intensity in order to allow a proper monitoring functionality; two different events reconstruction methods have been compared, showing the possibility to reconstruct the beam depth-dose profile and to retrieve the dose fall-off with millimeter precision.

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#### 1. Introduction

Ion beam therapy is a cancer treatment technique which is rapidly gaining importance in the global tumor therapy panorama. In addition to the already operational 70 clinical facilities, for more then 175000 patients already treated by the end of 2017, several new centers have been designed and approved for construction worldwide [Particle Therapy Cooperative Group, 2017]. The favorable feature of this treatment technique is connected to the peculiar energy deposition profile of charged particles as a function of depth in matter. As first observed by Bragg [Bragg and Kleeman, 1904], the depth-dose profile of charged particles shows a maximum close to the end of their range in matter; in addition to this, an enhanced biological effectiveness (capability of inducing DNA damages at fixed dose) has been verified for particles heavier than protons in several studies [Elsässer et al., , Weyrather et al., 1999].

The Bragg peak position corresponds to the maximum of the dose deposited in the patient and must be tuned to cover the target volume and, at the same time, spare the surrounding healthy tissues. In human body, the uncertainties in the material composition determination strongly limit the exploitation of the high ballistic precision of ion beams. The tumor volume is defined via CT scan in the standard clinical routine, and the treatment is planned via treatment planning software. In addition to the already cited uncertainties linked to the tissue composition, it must be noticed that other sources of treatment uncertainties are CT units conversion to ion stopping power, patient mispositioning, organ motion or morphological changes between treatment fractions (a standard treatment is divided into several fractions over several weeks, according to the tumor and patient characteristics). These uncertainties force the clinicians to fix relatively large safety margins around the planned treatment volume, up to 3.5\% + 3 mm [Paganetti, 2012]. The research and clinics community agrees on the fact that ion-range verification is one of the conditions (at present the main condition) required for a broader usage of ion beam therapy and for its further development. With the goal of fully exploit the ion beam therapy dosimetric potential, the monitoring should be in real-time and ideally in 3 dimensions, in order to be able to interrupt the treatment in case of severe issues, corresponding to important differences between the planned and delivered dose to the target volume or to surrounding organs at risk.

Several techniques have been considered worldwide for twenty years. Most of the studied techniques relies on the detection of secondary radiations generated during the slowing down process of incident ions, in particular during nuclear reactions. Among theses secondary radiations, positron emitters have been deeply studied in order to exploit positron emission tomography (PET) machines for treatment monitoring. PET techniques are based on the detection of the tow back-to-back 511 keV photons produced by the annihilation of positrons (emitted by the emitter fragments of nuclear reactions) with patient electrons, resulting in a delayed radiation which should be detected with time coincidences, allowing for an intrinsic background reduction. Nevertheless, the

monitoring with positron emitters secondary signal must deal with the mismatch between emitter production and positron production and annihilation. Even if the only available and functional range monitoring system in a clinical center is based on this technique [Enghardt et al., 2004], several clinical experience with commercial or adapted PET system already shown intrinsic limitations mainly connected to the ring geometry (not directly applicable to the treatment monitoring due to the presence of the beam) or in general to geometrical constraints limiting the field of view and the resulting system global efficiency and spatial accuracy (the limited detection angle generates artifacts in the final image) [Parodi, 2016]. The research is ongoing and new results are expected for the next years thanks to the introductions of new systems with adapted geometries, to the improvements in acquisition and reconstruction techniques and to the clinical introduction of time-of-flight systems, intrinsically able to improve the detector spatial resolution via interaction time information, and depth-of-interaction reconstruction, which will allow for a more precise spatial reconstruction for reduced angular artifacts effects.

In addition to positron annihilation products, a different kind of photons are emitted during ion irradiation. They emerge from the relaxation of excited nuclei in a wide energy range, between some hundreds of keV till about 8-10 MeV. After the first idea proposal published in 2003 [Stichelbaut and Jongen, 2003], these secondary products of particle treatment have been deeply investigated and the correlation of this gamma radiation to the ion depth-dose profile has been confirmed by several research groups, starting from [Min et al., 2006] for protons and [Testa et al., 2008] for heavier ions (mainly carbon). The so called prompt gamma-rays (PG) have the advantage to be emitted almost instantaneously after the beam interaction in the tissue, making them more adapted then PET 511 keV gammas for realtime monitoring. Consequently, different techniques have been proposed to exploit this signal for treatment monitoring purpose, with the related detection systems. Some methods are based on PG timing [Golnik et al., 2014, Krimmer et al., 2017] or energy [Verburg and Seco, 2014] information and rely on non-collimated systems; more complex detection apparatus can achieve an actual PG imaging, by exploiting mechanical or electronic collimation (i.e. with collimated gamma camera or Compton camera, respectively) for the photon selection (see e.g. [Min et al., 2006, Bom et al., 2012, Smeets et al., 2012, Roellinghoff et al., 2014, Priegnitz et al., 2015, Frances et al., 2010, Llosá et al., 2012, Kormoll et al., 2011, McCleskey et al., 2015, Matsuoka, 2015, Peterson et al., 2010, Solevi et al., 2016, Aldawood et al., 2017). Originally designed for astrophysics applications, the potential of Compton cameras

for medical imaging has been soon recognized [Todd et al., 1974] and then directly translated to the ion beam therapy monitoring domain. Such a gamma detection system is generally composed of two sections: a scatterer and an absorber. The scatterer is dedicated to the gamma Compton scattering, and should be so adapted in order to optimize the Compton scattering probability in the prompt gamma energy range, while reducing the so called Doppler broadening effect due to electron bounding

and motion [Ordonez et al., 1997]; this leads, in most of the cases, to the choice of a light material (low Z), segmented in several subsections. The absorber is devoted to the final absorption of the Compton scattered photons via photoelectric effect; it is often composed of segmented high-Z scintillating material. Slightly different Compton camera configuration can also achieve Compton electron tracking in the scattering detector [Frandes et al., 2010, Yoshihara et al., 2017], which results in additional information for the further reconstruction algorithm.

The collected interaction positions and energy depositions in the two detector sections are used to limit the emission point on the surface of a cone (or to a segment in the cone if the Compton electron track information is retrieved), via Compton kinematics reported in equation 1.

$$\cos\theta = 1 - \frac{m_e c^2 E_1}{E_2 (E_1 + E_2)},\tag{1}$$

where  $m_e c^2 = 511$  keV,  $E_1$  and  $E_2$  are the energies, respectively, deposited in the scatterer and the absorber. Once the cones are calculated, analytical or iterative algorithms are used to create the image of the prompt gamma emission distribution, with intrinsic 3 dimensional capability [McKisson et al., 1994, Kuchment and Terzioglu, 2016].

Several sources of incertitude and signal background are connected to this detection method. Considering a gamma interacting twice in the detector, i.e. once in the scatterer and once in the absorber, a complete photon energy absorption is needed since the initial photon energy  $(E_1)$  is not known a priori and the reported formula assumes valid the relation in equation 2.

$$E_0 = E_1 + E_2 (2)$$

An underestimation of the total initial energy (caused by a photon non-complete absorption in the absorber section or by the Compton electron escape from the scatterer section), leads to an underestimation of the Compton angle, so to a Compton cone reconstruction incertitude. For a three-interaction operating mode (making use of double scattered photons), the initial photon energy can be calculated analytically so that a complete absorption is not mandatory. In addition to this, the Compton kinematics formula does not take into account the initial bounding configuration of the Compton scattering electron, which creates a blur in the Compton angle reconstruction, resulting in the already cited Doppler broadening effect [Ordonez et al., 1997]. Furthermore, the time structure of the incoming particles plays an important role due to the detection principle, which is based on time coincidences between the two detector sections. The final image accuracy then suffers from false coincidences, generated by two prompt gammas interacting in the same time window or by the contamination of different kind of secondaries, mainly neutrons and protons. It is clear that excellent detector time resolution can limit the amount of random coincidences, as well as different background rejection methods can be applied to select real coincidences. Energy selections can be applied to

the collected coincidences [Polf et al., 2009, Hilaire et al., 2016], or the homogeneous neutron background can be reduced via time-of-flight information [Testa et al., 2010]. A French collaboration of 4 research institutions (Institut de Physique Nucleaire and Centre de Recherche en Acquisition et Traitement de l'Image pour la SantÃl' in Lyon, Laboratoire de Physique Subatomique et Cosmologie in Grenoble, Centre de Physique des Particules in Marseille) is focused on the design and development of gamma detectors for online ion beam therapy monitoring. The project includes the development and clinical test of a Compton camera prototype based on semiconductor and scintillator detectors [Krimmer et al., 2015]; the feasibility of its clinical application is studied in this work with Monte Carlo simulations.

In a previous study by Ortega and colleagues [Ortega et al., 2015] a detailed analysis of the noise sources for Compton imaging in proton therapy monitoring is presented, and the clinical application of this method for detecting range shifts is tested. The simulation study, performed with a simple Compton camera prototype composed by multiple LaBr<sub>3</sub> layers with a ideal mono-energetic proton beam impinging on a PMMA phantom, showed the relative expected rate of prompt gammas and neutrons, and the resulting rate of random coincidences ranging from 19 % to more than 60 % depending on the beam energy and the coincidence time window. This amount of fake events leads to complex reconstruction scenarios, where the identification of a 3 mm range shift is not clear for all cases.

Starting from these results, the ClaRyS camera performance are here studied as a function of the gamma energy in the prompt gamma energy range, and the possible application of this detector as depth-dose profile monitor during ion beam therapy clinical treatment is analyzed by investigating the effect of the beam time structure on the rate of true and random coincidences, as well as on the background level. Clinical proton and carbon beams impinging on an homogeneous PMMA phantom are simulated to reproduce treatment conditions and analyze the prompt gamma detection resulting scenario. A beam intensity reduction with respect to treatment intensities is needed for monitoring purpose, with the aim of increasing the ratio between true and random coincidence events: the maximum beam rate tolerance is here estimated. Finally, the Compton camera precision in the identification of the dose profile fall-off is reported.

#### 2. Material and methods

#### 2.1. Simulation setup

The monitoring system modeled in this simulation work is a Compton camera prototype under development within the French collaboration CLaRyS (Contrôle en Ligne avec Rayonnements Secondaires) including five institutions: IPN Lyon, CREATIS Lyon, LPC Clermont-Ferrand, CPP Marseille and LPSC Grenoble. The detectors detailed characteristics can be found in [Krimmer et al., 2015]. A scheme of the simulation setup is given in figure 1.

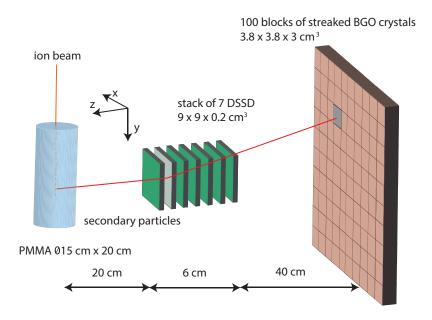


Figure 1: Scheme of the simulation setup: a PMMA cylindrical phantom is set in front of the Compton camera prototype. The Compton camera is composed of a stack of 7 DSSDs (scatterer) and a plan of 100 single BGO blocks. The set distances are realistic for clinical conditions. This geometrical configuration has been used for all the simulations presented in this work.

Like most of the Compton camera devices, the CLaRyS prototype includes a scatterer and an absorber. The scatterer consists of seven parallel planes of silicon detectors (double-sided silicon strip detectors, DSSDs),  $9 \times 9 \times 0.2$  cm<sup>3</sup>, with 1 cm distance between the centers of two neighboring planes, while the absorber is composed of an array of  $10 \times 10$  BGO blocks ( $3.5 \times 3.5 \times 3.0$  cm<sup>3</sup> each) placed at 40 cm from the last silicon layer (center-to-center distance). The ion beam interacts with a cylindrical PMMA† phantom (15 cm diameter and 20 cm length) placed in front of the Compton camera as target. It is placed 20 cm far from the first silicon plane (center-to-center distance) in order to fit with realistic clinical conditions.

The silicon detectors have a strip pitch of 1.4 mm, for a total of 64 strips per side (double-

sided readout based on electron and hole pairs collection). The strips are not reproduced in the simulation code, and the transverse spatial resolution is set to 0.9 mm FWHM at the reference energy of 1 MeV, according to preliminary measurements performed on smaller detector prototypes. Concerning the parallel direction, the interaction position is set to the center of the involved silicon plane.

Regarding the BGO blocks, their entrance surface is streaked in a  $8\times8$  matrix of pseudopixels, 4.4 mm² side, and the readout is performed via 4 photo multiplier tubes. The position reconstruction is achieved via Anger logic for the real detector, but a monoblock crystal is simulated here for simplicity. The events are selected to be limited to a single block component based on spatial analysis, and the interaction position is reconstructed via center of gravity calculation if multiple interaction occur. An incertitude contribution, randomly extracted by a gaussian of 5 mm FWHM, is added to the reconstructed position to fit with the geometrical features. For what concerns the parallel direction, given the fact that the employed BGO blocks have not depth of interaction reconstruction capabilities, the interaction position is fixed to the center of the mono-block crystal.

Preliminary characterization measurements allowed to estimate the energy resolution of the BGO blocks, which is accordingly set to 17% FWHM at the reference energy of 667 keV (a 137-cesium source has been used for the measurements). The energy resolution of the silicon detector is set to 2.3 keV (RMS) according to the design expectations; no characterization data are yet available for an instrumental estimate.

The time resolution has been set to 3.0 ns FWHM for the BGO blocks and to 15.0 ns FWHM for the silicon slabs, according to preliminary measurements performed on test detector modules at the GANIL (Grand Accelerateur National d'Ions Lourds) center in France.

The detector resolutions play an important role in the Compton camera performances. The absorber spatial resolution influences the position of the apex of the Compton cone, as well as its axis orientation. The scatterer energy resolution determines the Compton cone aperture angle. The time resolution impacts the coincidence window between the absorber and the scatterer, and so its true/random coincidence discrimination capabilities.

The CLaRyS project also includes the development of a beam tagging hodoscope, composed of scintillating fibers read out by multi-channel photomultipliers. This detector is used to synchronized the beam time and space structure to the prompt gamma detection in order to tune the detection window reducing the background contamination. This detector section is not included in the simulation, but its time resolution has to be taken into account for the time-of-flight discrimination. It is set to 1 ns FWHM.

The detectors spatial, energy and time resolutions are summarized in table 1.

The Monte Carlo simulation study is performed with the Geant4 toolkit, version 9.6

<sup>†</sup> PolyMethylMethAcrylate

<sup>†</sup> Bismuth Germanate - Bi<sub>1</sub>2GeO<sub>2</sub>0

Table 1: Estimations of reachable resolutions with the detectors. Those resolutions are applied during the simulations.

| Resolution (FWHM) at 1 MeV | Scatterer | Absorber | Hodoscope |
|----------------------------|-----------|----------|-----------|
| spatial [mm]               | 0.9       | 5        | /         |
| energy [%]                 | 2.3       | 17       | /         |
| ${f timing}  [{f ns}]$     | 15        | 3        | 1         |

patch 02. Geant4 has been developed at CERN (Counseil europÄl'en pour la recherche nuclÃl'aire) for high energy physics experiments, but it has been shown that it can be used for ion beam therapy studies [Cirrone et al., 2011, Toshito et al., 2010]. However, some improvements are still needed in order to extend the hadronic models to low energy applications [Dedes et al., 2014].

The particle interactions in matter are described in this work by means of different models, listed in table 2. Additionally, the Doppler broadening and the photon polarization effects are taken into account.

Table 2: Hadronic models used in the Geant4 simulations.

| Processus        | Protons         | Ions   | Neutrons                                  |  |  |
|------------------|-----------------|--|---|--|--|
| Electromagnetic  |                 | $\operatorname{standard}_{\operatorname{option}3}$ |   |  |  |
| Inelastic        | G4BinaryCascade | G4QMDReaction                                      | G4BinaryCascade                           |  |  |
|                  |                 | (G4IonsShenCrossSection)                           | + G4NeutronHPInelastic (<19 MeV)          |  |  |
| Elastic          | G4LElastic      | G4LElastic   | G4LElastic + G4NeutronHPElastic (<19 MeV) |  |  |
| Fission          | /               | /  | G4LFission + G4NeutronHPFission(<19 MeV)  |  |  |
| Capture          | /               | /  | G4LCapture + G4NeutronHPCapture (<19 MeV) |  |  |
| Radioactivedecay | /               | G4Radioactivedecay                                 | /   |  |  |

The two main beam particles used in clinics are considered in this work: protons and carbon ions. The beam range of interest is 15.2 cm in the PMMA phantom, and the associated energy is 160 MeV for protons and 305 MeV/n for carbon ions. In order to reproduce a realistic clinical beam, the beam transverse dimension is modeled with a gaussian distribution with a standard deviation of 5 mm for protons at 160 MeV and 3.5 mm for carbon ions at 305 MeV/n. The Compton camera setup does not change for the different incident particles. The beam intensity for a spot in pencil beam scanning (PBS) mode for protons is 10<sup>8</sup> particles and 10<sup>5</sup> particles for carbon ions.

The beam time structure is applied in the data analysis stage, described below.

#### 2.2. Data treatment

2.2.1. Time structure and treatment Two different beam time structures have been considered for this study, related to two kinds of accelerators used in clinical practice: the

IBA C230 cyclotron for protons (used in 16 clinical centers worldwide) and the Heidelberg (Germany) synchrotron installed in the Ion Therapy Center (HIT) for carbon ions. Even if the beam microstructure changes depending on the ion energy and the beam intensity, the microstructure is modeled according to the reported features at a specific energy and intensity for simplicity. For protons at 160 MeV, the primary particles are grouped in bunches of 2 ns at a frequency of 106 MHz (9.42 ns) [F Roellinghoff, 2014]. The clinical beam intensity is 3.2 nA which corresponds to about 200 protons per bunch. Concerning the carbon ion beam at 305 MeV/u, the estimated microstructure is composed of 30 ns bunches at a frequency of 5.9 MHz (170 ns). The clinical beam intensity for carbon ions is  $5 \times 10^7$  ions/s, corresponding to about 9 ions per bunch. This beam structure is extrapolated from measurements performed by our group in 2013 at HIT; the beam time structure was measured for 200 MeV/u and 400 MeV/u primary ion energy with a two-fiber hodoscope (basic prototype of the one at present under development) and the spill signal was given by the accelerator. Figure 2 shows the beam structure for carbon ions at 400 MeV/u. The pulses have a spill period of 150.2 ns and each bunch is approximately 21.5 ns. The mentioned measurements have shown that the spill phase changes during the extraction: this implies that the HF signal from the synchrotron can not be used to locate the pulses, so that the use of the hodoscope seems required for time-of-flight background rejection purposes.

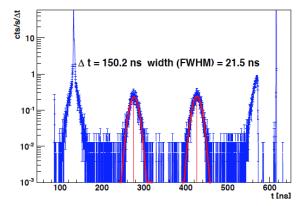


Figure 2: Time structure measured from a carbon ion beam at 400 MeV/u delivered at HIT. The pulses have an extraction period of 150.2 ns and the bunches are 21.5 ns FWHM. The measurement was achieved with a two-scintillating-fiber hodoscope.

The coincidence window (between scatterer and absorber events) is set to 40 ns, centered on each absorber detected interaction. This value is adapted to the detectors time resolutions.

Table 3 resumes the presented beam time structures and coincidence reconstruction features.

Table 3: Description of the two beam structures studied: the IBA cyclotron C230 for protons and the synchrotron installed at the Heidelberg Ion Therapy Center (HIT) in Germany for carbon ions. The beam structures are applied to the simulation data.

|                   |                         | Protons                           | Carbon ions                    |
|-------------------|-------------------------|-----------------------------------|--------------------------------|
| Clinical features | Facility                | IBA Cyclotron C230                | Synchrotron at HIT             |
|                   | Clinical intensity      | $2 \times 10^{10} \mathrm{\ p/s}$ | $5 \times 10^7 \text{ ions/s}$ |
|                   | Energy                  | $160~{ m MeV}$                    | $305~{ m MeV/u}$               |
| Beam structure    | Bunch time [ns]         | 3.2                               | 30                             |
|                   | Period [ns]             | 9.4                               | 170                            |
|                   | Primaries /bunch        | 217                               | 9                              |
| Detectors         | Coincidence window [ns] | 40                                | 40                             |
|                   | Time resolution [ns]    | Si: 15 and BGO: 3                 |                                |

2.2.2. Data selection: time and energy cuts As already explained in the introduction, the Compton detection principle is based on a double interaction in the scatterer and absorber section, where an interaction is defined as an energy deposit in a detector module. As discussed in section 2.2.1, the coincidence reconstruction relies on a defined time window, fixed according to the involved detector resolution. In a simulation environment, different kinds of coincidences events can be distinguished and studied:

- real coincidences: created by a single photon first interaction in at least one scatterer plane and then in a single absorber block;
- quasi-simultaneous interaction from two secondary particles;
- double interaction from the same particle different from a gamma.

In a real clinical environment there is no way to select the real coincidences, so that the collected data are affected by a certain number of the so called random coincidences. The amount of random coincidences depends on the detector time resolutions, on the fixed time coincidence window and on the beam time structure, for fixed beam energy, phantom composition and camera prototype setup.

In figure 3 a schematic view of the different kind of coincidences is presented.

In addition to this, the prompt gamma measurement is contaminated by other secondary particles produced by the beam interaction with the patient/phantom, mainly massive and charged particles (protons, neutrons). As already mentioned, it has been demonstrated in previous work by our group that a time-of-flight discrimination is possible and effective in reducing this source of background. In fact, the photons are moving at the speed of light while the massive particles approach the detector at a lowest speed. The time information coming from the hodoscope and from the absorber can be then combined to define fixed detection time window and reject all the events outside the window (3). In this work, the time between the incident particle creation and the secondary particle detection in the absorber is considered as the time-of-flight. The hodoscope time resolution  $u_{hodoscope}$  (1 ns FWHM) is applied to the primary particle creation time, with a contribution randomply extracted from a gaussian with  $\sigma = 1/2.35$  ns, while the

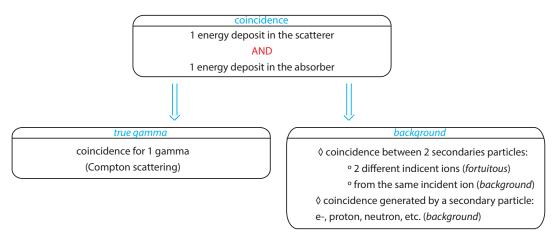


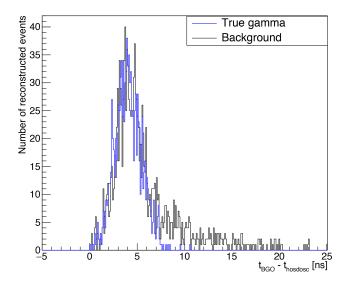
Figure 3: Diagram showing the different definitions of coincidences in the Compton camera.

detection time in the absorber is affected by the absorber time resolution.

$$TOF_{theoretical} = t_{absorber} - t_{hodoscope} TOF_{simulation} = t_{absorber} + t_{creation} + u_{hodoscope}(3)$$

The time-of-flight spectrum resulting from the simulation shows that the coincidences of interest (produced by prompt-gamma rays) are included in a window between 0 and 6 ns (figure 4). Therefore, all the coincidences with a TOF higher than 6 ns have been rejected, reproducing a possible clinical background rejection scenario.

Figure 4: Time of flight spectrum obtained by means of the simulation for a proton beam at 160 MeV and  $1 \times 10^8$  incident protons. The detection time for the absorder is given by the tag for the first deposit energy. The blue curve represents the time of flight for true events and the black one represents the background.



In addition to the time-of-flight based selection, energy thresholds are also defined for the event detection. On a single detector basis, 50 keV is set as lower threshold for the silicon layers and 100 keV for the absorber. On a complete event basis, a total absorbed energy lower limit is set to 1 MeV.

### 2.3. Reconstruction algorithm

Once the coincidences are defined and selected according to the fixed physical cuts, the prompt-gamma emission point has to be reconstructed for each event. This can be done via analytic or iterative algorithms based on the Compton kinematics (see section 1), presented in the following sections.

2.3.1. Line-cone algorithm The reconstruction via line-cone algorithm exploits the energy deposit and position information collected by the camera in addition to the beam path information. Thanks to the deposited energies in the detectors and the interaction positions, a cone surface is analytically defined via the Compton equation 1. We assume that the initial energy of the gamma ray is fully absorbed in the absorber. This hypothesis, combined to the detector energy resolutions, lead to a potential uncertainty on the cone aperture. The interaction position in the scatter gives the cone apex and the line connecting the interaction position in scatterer and absorber gives the cone axis. In order to simplify the reconstruction, the beam direction is used to limit the possible solutions (laying on the reconstructed cone surface) to two points (intersection of the beam direction and the reconstructed cone). The set of all the reconstructed points gives the emission source distribution. The obtained final image is then the 2 dimensional projection of the prompt gamma emission spectrum.

2.3.2. LM-MLEM algorithm The iterative methods allow to get a 3D image reconstruction, potentially by taking into account the spatial resolution and the energy resolution of the detectors. Few iterative algorithms have been developed for Compton event reconstruction. [Schone et al., 2010, Zoglauer et al., 2011, Gillam et al., 2011, Mackin et al., 2012, Lojacono et al., 2013].

The List-Mode Maximum Likelihood Expectation Maximization (LM-MLEM) algorithm is a MLEM version which allows to free from sinograms and to reconstruct the image directly from the list of detected events. The first step is to define the volume which includes the origin of the prompt gamma ray detected. This volume is divided into equal voxels and the emission intensity is assumed homogeneous for each voxel j, with a Poisson distribution of parameter  $\lambda_j$  (a vector of the emissions intensities of all the voxels). The algorithm is based on a system matrix T composed of the coefficients  $t_{ij}$  which represents the probability that a photon produced in the voxel j is detected in coincidence by the Compton camera as an event i. The probability for a gamma detected in coincidence to be emitted from the voxel j is denoted as  $s_j$ . The LM-MLEM

algorithm starts with an initial value  $\lambda^{(0)}$ , which can be the simple back-projection reconstruction. The iterations rely on the following recurrence relation:

$$\lambda_j^{(l+1)} = \frac{\lambda_j^{(l)}}{s_j} \sum_{i=1}^{N_{\gamma}} t_{ij} \frac{1}{P_i^{(l)}}, \quad \text{avec} \quad P_i^{(l)} = \sum_{k=1}^{N_{\nu}} t_{ij} \lambda_k^{(l)}, \tag{4}$$

where  $N_{\gamma}$  is the number of detected events and  $N_v$  is the number of voxels in the image.

The LM-MLEM algorithm used for this study is the one developed by the CREATIS research group in Lyon [Maxim et al., 2009, Lojacono et al., 2013, Maxim, 2014, Hilaire et al., 2014].

For each photon detected, the system matrix T is calculated by taking into account the uncertainties on the angle between the source and the involved scatterer plane and the angle between the scatterer plane and the absorber involved module. The matrix elements  $t_{ij}$  are calculated as:

$$t_{ij} = K(\beta_i, E_{tot}) \frac{|\cos(\theta_{\overrightarrow{V2V1}})|}{V_2 V_1^2} \int_{M \in v_i} \frac{|\cos(\theta_{\overrightarrow{V_1M}})|}{V_1 M^2} h_i(M) dv, \tag{5}$$

where  $\beta_i$  is the Compton scattering angle,  $V_1$  the interaction position in the scatterer,  $V_2$  the interaction position in the absorber,  $h_i$  the spatial kernel which models the uncertainties on the Compton angle for each voxel M,  $K(\beta_i, E_{tot})$  the differential cross section and v the reconstructed volume.

In order to simplify and speed up the calculation of the  $t_{ij}$  matrix, the voxels located far from the reconstructed cone are set to 0. The distance between the cone and the voxel is calculated by taking the voxel center as reference point. The spatial resolutions are not included in this version of the algorithm.

For each iteration, the matrix T is stored and the reconstructed image can be produced via a Matlab simple interface.

#### 2.4. Precision estimation

The camera precision is defined as the difference between the expected Bragg peak position (according to the treatment planning) and the Bragg peak position reconstructed through the data. This is a critical parameter which will be used to fix a decision threshold during a ion beam treatment.

In this study a reference profile has been defined as the reconstructed emission vertex profile at high statistics, via analytic (line cone) and an iterative (LM-MLEM) algorithms; this will be used to mimic the treatment planning. This Bragg peak position is compared to the ones reconstructed at lower statistics (close to clinical conditions) via analytic and iterative algorithms.

The high statistic profile has been obtained with  $2 \times 10^{10}$  incident protons (figure 5a) for

the line-cone reconstruction and with  $1 \times 10^{10}$  incident protons (figure 5b) for the LM-MLEM reconstruction method. The SmoothKern method, with the Nadaraya-Watson regression, is used to smooth the reference profiles in order to reduce relative statistic fluctuations.

A region of interest (ROI) ranging from y = 0 mm to y = +100 mm is defined around the expected Bragg peak position, located at y = +50 mm in the phantom. The reference profiles are modelled in the ROI by a linear combination named Non-Uniform Rational Basis Splines (NURBS).

A new independent profile should be simulated at low statistics for the precision calculation. However, with the aim to speed up the analysis, a random extraction from the NURBS profile (5c and 5d) is performed following the Poisson law, in order to select reduced data sets. The desired low statistics ranges from  $10^8$  to  $5 \times 10^9$  incident protons.

The minimal distance between the NURBS reference profile and the low statistic reconstructed profiles is estimated thanks to the  $\chi^2$  method. For the  $\chi^2$  estimate, the low statistics profile is defined around the fall-off position from y=+30 mm to y=+70 mm. The minimization process tests the profiles at low statistics on a 60 mm variation range, between -30 mm and +30 mm with respect to the initial position, with a step of 0.1 mm. The  $\chi^2$  is calculated as follows:

$$\chi^2 = \sum_{i=1} \left( y_{sample,i} - y_{NURBS,i} \right)^2, \tag{6}$$

where  $y_{sample}$  is the number of coincidences for the low statistics profile,  $y_{NURBS}$  is the number of coincidences for the reference profile NURBS (scaled at the same low statistic) and i the step number.

The global minimum of all the reconstructed profiles is then retrieved. The figures 5e and 5f show the distribution of  $\chi^2$  calculated for a low statistic profile at  $10^8$  incident protons.

A total of a thousand profiles at the low statistic are generated (named realizations) and the  $\chi^2$  minimization is applied to each of them. The standard deviation of the distribution resulting of the thousand results gives the precision of the camera for a given number of incident protons. The figure 5g and 5h show the distributions at  $10^8$  incident protons for the line-cone and LM-MLEM algorithms.

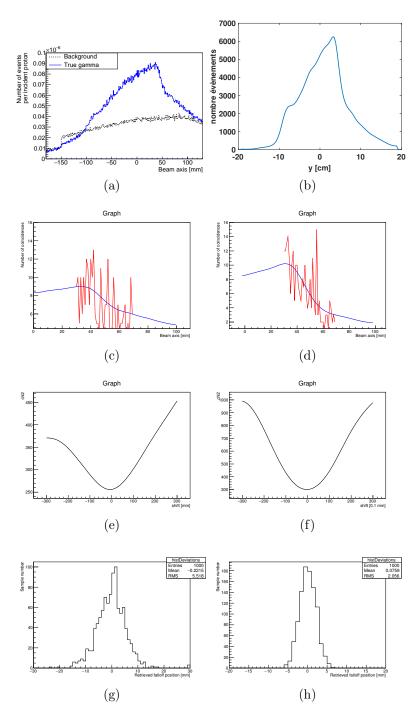


Figure 5: Treatment comparaison for the same proton simulation with the line cone algorithm (left column: figures 5a, 5c, 5e, 5g) and the LM-MLEM algorithm (right column: figures 5b, 5d, 5f, 5h). The first row give the reconstructed profils for  $2 \times 10^{10}$  incident protons (figure 5a) and for  $1 \times 10^{10}$  incident protons (figure 5b) respectively. The figures 5c and 5d are showing the reference curve (blue) and the estimated curve with Poisson's law at  $1 \times 10^8$  incident protons. The figures 5e and 5f are the  $\chi^2$  minimization results for one realization. The minimum of the curve gives the best fit between the reference curve and the low statistics one. Finally, the figures 5g and 5h represents the results for  $1000 \chi^2$  minimizations. The Compton camera precision is estimated thanks to the standard deviation of the distribution.

#### 3. Results

### 3.1. CC efficiency

The absolute efficiency is crucial for the Compton camera performances. It affects directly the number of detected events by the camera and so the reconstruction quality of those events. The absolute efficiency  $\epsilon$  is defined as:

$$\epsilon = \frac{N\gamma_{recons}}{N\gamma_{total}},\tag{7}$$

with  $N\gamma_{recons}$  the number of gamma events in coincidences,

 $N\gamma_{total}$  the total number of gamma emitted: 10<sup>8</sup>.

The absolute efficiency is presented in function of the localization of a monoenegertic gamma point source in comparison to the Compton camera center. The setup is the same as figure 1, but the point source is in the air and not in a PMMA phantom. The point source is moved from -300 mm to +300 mm with a step of 20 mm following the transverse axis of the camera (axis y) and the gamma source is monoenergetic [300 keV, 6 MeV].

The figure ?? shows the absolute Compton camera efficiency in function of the gamma source position. The figure ?? gives the results without any cut in energy on the detectors and the figure ?? is with a cut applied on the energy deposited in the detectors: 50 keV for the scatterer and 100 keV for the absorber.

The absolute efficiency is between  $4 \times 10^{-4}$  at 300 keV and  $1.5 \times 10^{-4}$  at 6 MeV (figure ??). The efficiency is higher for low gamma energies because the probability of interaction in the absorber is decreasing with the energy. Moreover, when the point source is far from the camera center, the efficiency drops quickly. This effect is marked for high energies. In fact, the incident gamma will be less deflected in the scatterer for the same energy deposited compared to a low energy gamma. The incident gamma of interest in hadrontherapy are around 1 MeV, so it is important to well positioned the Compton camera to optimise the efficiency.

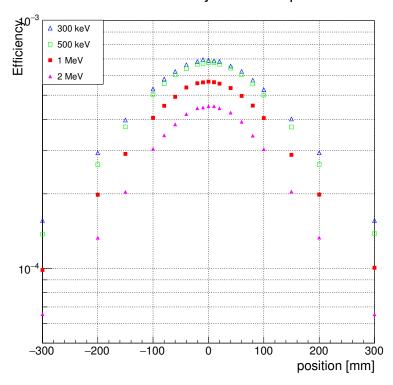
Another aspect to taking into account is the energy cut applied to the detectors (figure ??). The detectors are not perfect and cuts are mandatory to eliminate the electronic noise. If a gamma has a small Compton scattering angle, the energy deposited in the scatterer will be small. That is why, the camera efficiency decreases when the gamma source is centred with the camera. The limited energy cut is the one applied to the scatterer. This point is important to notice for an application of the Compton camera in nuclear medicine.

#### 3.2. Beam intensity

The beam intensity plays an important role for the Compton camera concerning its capability to distinct events in coincidence. In the simulation, the beam intensity is modeled by an average number of particles per bunch. The exact number of particles

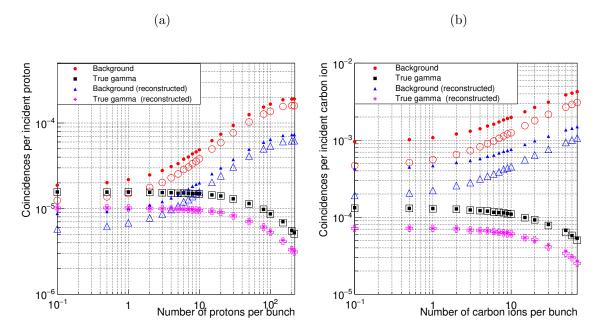
Figure 6: Absolute Compton camera efficiency function of the gamma source position. The figure ?? shows the camera efficiency with no energy cut applied. It models the perfect Compton camera. The figure ?? takes into account cuts in energy: 50 keV for the scatterer and 100 keV for the absorber. The value of the energy cut can be different for the final prototype depending of the detector energy resolutions reach. The analysis is done for energies from 300 keV to 6 MeV.

# Detection efficiency vs source position



in each bunch is given by a random draw in a Poisson distribution, where the mean value is the beam intensity chosen. The range of intensities was chosen in order to cover almost all the possibilities: from a very low beam intensity to the clinical beam intensity. Therefore, for proton and carbon ion, the lowest beam intensity is set to 0.1 particles per bunch in average and for protons goes up to 217 protons per bunch when in case of carbon ion, it goes up to 70 particles per bunch. The coincidence yields are scaled per ion incidents and the beam intensity per average ions per bunch. The true coincidences represent a coincidence in the camera by the same gamma ray. The background corresponds to all the other coincidence types. The simulations are done for a statistic of  $10^8$  in case of protons and  $2 \times 10^5$  for carbon ions. Those statistics correspond to the number of ions used in clinics to treat a spot.

Figure 7: Coincidences yield for protons (figure ??) and carbon ions (figure ??) in function of the beam intensity. The intensity is given for a number of incident particles per bunch. The distinction between the filled markers and the empty ones are the time of flight discrimination applied in the case of empty markers. Moreover, the yields are given before and after the profile reconstruction with the line cone algorithm.



At the clinical beam intensity, the high background level is mainly due to the random coincidences. In fact, the probability to detect two radiation coming from two different incident particles increases with the number of incident particles per bunch. Another issue is the single rate of events detected by each detector at those high intensities. For instance, at the clinic beam intensity in proton therapy, the single rate on the absorber is around 300 MHz and on the first silicon layer it is around 20 MHz. The current electronic front end and acquisition system are not able to treat this amount of data coming from all the detectors. As a result, it appears that it is impossible to use the Compton camera at a clinical beam intensity for the treatment monitoring in ion therapy.

Nevertheless, if the intensity decreases enough to avoid almost all the random coincidences, the monitoring seems more feasible. In addition, it can be supposed that the time of flight discrimination will improve the signal to the background ratio at low intensities by suppressing the coincidences induced by charged particles. Indeed, the charged particles are slower than gamma rays which move at the light speed.

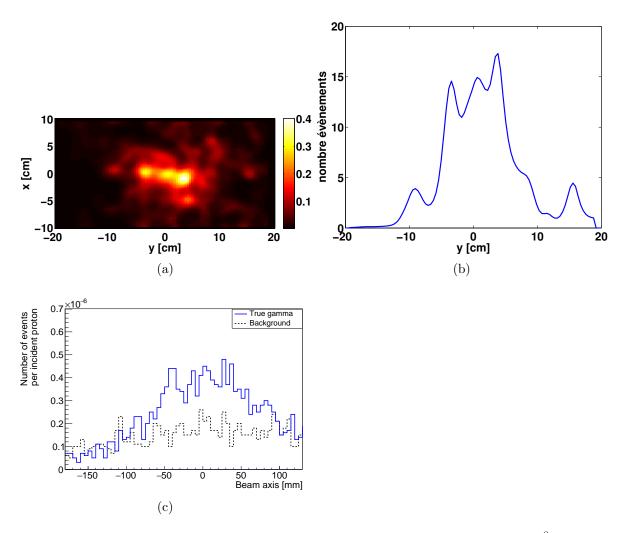


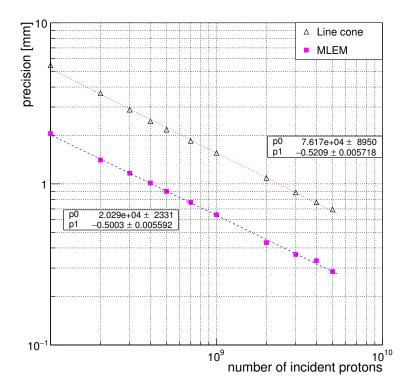
Figure 8: LM-MLEM reconstruction for a proton beam at 160 MeV and  $10^8$  incident protons. The events are detected in coincidence and the beam intensity is reduced to 1 proton per bunch. The PMMA target has a diameter of 15 cm and is 20 cm long. The Compton camera is centred at y = +50mm. The time of flight discrimination is applied and 20 iterations are realized. The figure ?? represents the reconstruction in 2D for the plan (x,y). The position x = 0mm is the center of the PMMA phantom and y axis the direction of the proton beam. The figure ?? shows the profil corresponding to the figure ?? and following the axis y. The Bragg peak is localized at y = +50mm. The figure ?? is the profil obtained by means of line cone algorithm for the same simulation datas.

### 3.3. Comparaison LM-MLEM vs Line cone reconstruction

#### 3.4. Compton camera precision

The information given by a control device as the Compton camera has to be viable and as precise as possible. In order to estimate the precision of the Compton camera, a method is used.

Figure 9: The Compton camera precision is estimated with two different algorithms: line cone and LM-MLEM. The precision is given for  $1 \times 10^8$  to  $5 \times 10^9$  incident protons. A linear fit is realized in order to obtain the slope of the results (p1 parameter). The graphic is scaled in log-log.



# 4. Discussion

DISCUSSION

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