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Additional Information



Novel Method to Measure the Intrinsic Spatial Resolution in PET Detectors Based on Monolithic Crystals

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Abstract

The main aim of this work is to provide a method to retrieve the intrinsic spatial resolution of a gamma-ray detector block based on monolithic crystals within an assembled scanner. This method consists on a discrimination of the data using a software collimation process. The results are compared with an alternative method of separating two detector blocks far enough to produce a "virtual" source collimation due to the geometric constraints on the allowed coincidence event angles.

A theoretical model has been deduced to fit the measured light distribution profiles, allowing estimating the detector intrinsic spatial resolution. The detector intrinsic spatial resolution is expected to follow a Gaussian distribution and the positron-emitter source shape, given the small size of a ²²Na source with 0.25 mm in diameter, can be assumed to follow a Lorentzian profile. However, the collimation of the data modifies the source shape that is no longer a pure Lorentzian distribution. Therefore, the model is based on the convolution of a Gaussian shaped distribution (contribution of the detector) and a modified Lorentzian distribution (contribution of the collimated source profile) that takes into account the collimation effect.

Three LYSO crystals geometries have been studied in the present work, namely a 10 mm thick trapezoidal monolithic block, and two rectangular monolithic blocks with thicknesses of 15 mm and 20 mm, respectively. All the blocks have size dimensions of 50 mm \times 50 mm. The experimental results yielded an intrinsic detector spatial resolution of 0.64 ± 0.02 mm, 0.82 ± 0.02 and 1.07 ± 0.03 mm, for the 10 mm, 15 mm and 20 mm thick blocks, respectively, when the source was placed at the center of the detector. The detector intrinsic spatial resolution was moreover evaluated across one of the axis of each crystal. These values worsen to an average value of 0.68 ± 0.04 mm, 0.90 ± 0.14 and 1.29 ± 0.19 mm, respectively, when the whole crystal size is considered, as expected. These tests show an accurate method to determine the intrinsic spatial resolution of monolithic-based detector blocks, once assembled in the PET system.

Keywords: Gamma ray detectors, Positron Emission Tomography, Intrinsic spatial resolution, SiPM array, Monolithic blocks

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1. Introduction

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source must be taken into account.

In the field of Nuclear Medicine, Positron Emission Tomography (PET) scanners are widely used. They provide quantitative functional information of the subject under study. In order to record the 511 keV gamma rays resulting from the positron-electron annihilation, these scanners have been typically based on scintillation crystal arrays [1]. However, in recent years, the interest of using PET detector blocks based on monolithic crystals has increased. This interest is reflected not only from the academic point of view [2]-[6], but also from the commercial one [7]-[9]. The main advantages of continuous crystals, such as the characterization of the light distribution [10][11], the estimation of the gamma ray impact coordinates [12]-[14], or its spatial, energy and time resolution capabilities [15]-[19], have been widely described elsewhere. One of the most important benefits concerning the use of continuous blocks, in contrast to pixelated crystals, is the possibility to estimate the photon depth of interaction (DOI) coordinate without the need of extra photosensors or scintillator material [20]-[25]. The intrinsic spatial resolution of detector blocks based on crystal arrays is typically given by the system capabilities to resolve all crystal elements [21][22]. Also, it is related to the signal-to-noise ratio, provided by the peak-to-valley ratio in flood images resolving these pixel elements. However, when monolithic crystals are used, accurate determination of the intrinsic spatial resolution is challenging. In those crystals, the detector spatial resolution is limited by several factors such as detector size, photosensor type and distribution, photon DOI and scatter events, to name but a few. These factors cannot easily be isolated and studied experimentally. For this reason several works have focused on exploring the resolution limits through Monte Carlo simulations or other models [26]-[28]. Other factors, namely the non-collinearity of the annihilated photons or the positron annihilation range, contribute to the reconstructed image resolution and are not taken into account when referring to the detector itself. However, it should be pointed out that if the intrinsic detector spatial resolution is determined by using positron emitters, then in addition to the source size, the positron annihilation range of the

In detectors based on continuous crystals, it has been suggested obtaining the intrinsic spatial resolution by having a reference detector at a far distance generating a narrow gamma-ray beam. This technique has been used not only in the PET field [16], but also in other areas of research such as Astrophysics [29]. However, this is not easily applicable for two main reasons. First, long separation distances are required and, therefore, in order to have enough statistics, high activity sources or long measuring times are needed. Second, this is impracticable if the intrinsic detector resolution is aimed to be obtained once the PET scanner is assembled. Other methods are based on using physical collimators (tungsten or lead) with small apertures. However, using collimators increases the measurement time and produces a magnification effect of the source profile at the entrance face of the scintillator block (proportional to the collimator thickness) that has to be taken into account when calculating the detector intrinsic resolution. The aim of this work is twofold. We demonstrate a method to measure the detector intrinsic spatial resolution of a PET system when the scanner is already assembled. The method we are proposing is based on software collimation of the data, thus, avoiding the need of moving away one of the detectors. In the following, we will refer to this method as "software collimation". We have compared the proposed method against the aforementioned "distant detector" approach. The second topic is about modelling the expected measured resolution profile. This model includes the essential processes contributing to the shape of the measured resolution profile. When a fitted model has converged to the measured resolution profile, the FWHM of the intrinsic resolution PDF can be extracted.

2. Materials

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2.1. Monolithic scintillators

The monolithic crystal blocks under study are composed of LYSO material. Three different detector block geometries have been tested, namely a trapezoidal (50 mm × 50 mm exit face, 47.9 mm × 47.9 mm entrance face) block with a thickness of 10 mm [5], and two rectangular blocks (50 mm × 50 mm base) with thicknesses of 15

mm [30], and 20 mm [23], see Fig. 1 left, center and right, respectively. For all the scintillation blocks, all faces are polished and the lateral ones black painted. The entrance face is coupled to a retro-reflector (RR) layer [23]. The RR layer is composed of several corner-cubes structures that are made of three-sided prisms (mutually perpendicular) and reflect light back directly towards the source, displaced a small distance (120 µm), but preserving the light distribution shape. The use of these devices has a positive impact on the spatial resolution as demonstrated in [23][30][31]. For a given LYSO geometry, two identical detector blocks have been used in coincidence mode, one of them working as the reference detector.



Fig. 1. Photograph of the three LYSO monolithic blocks used for the experiments. Left, trapezoidal block 10 mm thick. Center, rectangular block 15 mm thick. Right, rectangular block 20 mm thick. The retroreflector layer coupled to the entrance face of the blocks can be seen through the crystals.

2.2. Photosensor and readout

Custom photosensor arrays composed by 16×16 SiPMs (Silicon Photomultipliers) with 3×3 mm² active area each (SensL, Cork, Ireland), and a pitch of 3.26 mm, were used. The SiPM matrix is operated at a bias voltage of 29 V, 4.5 V over the breakdown voltage [1]. Each SiPM has been connected to a readout circuit that provides outputs for each row and column of the photosensor matrix [32]. The row and column readout electronics allow one to characterize the light distribution (LD) originated from the interaction of a gamma ray within the monolithic scintillator. The projection of the LD onto the X and Y axes makes it possible to determine the planar X and Y coordinates, and also the photon DOI (Z coordinate).

2.3. Radioactive source

- 2 The ²²Na radioactive source used across all the experiments has a spherical shape with 0.25 mm in diameter
- 3 and placed in the center of a PMMA encapsulation with 6 mm height and 25.4 mm in diameter. The activity of
- 4 the source was low, about 7.4 kBq and, therefore, pile-up events due to high rates are not expected.

3. Methods

The readout electronics of each detector module provides digitized information of the LD projections for both X and Y axes [1][5][23][30]. In order to improve the calculation of the X and Y impact positions near the crystal edges, the centroids of these distributions are calculated through the center of gravity algorithm rose to a power of 2 [33]. The photon impact DOI is estimated by the ratio of the sum of all 16 signals (photon energy, E) to the maximum signal value (E/I_{max})_{r,c} for row and columns (r,c) [23]. The final DOI value is assigned to the mean value of (E/I_{max}) obtained for r and c.

For all measurements the detector blocks temperature was monitored using sensors on the rear part of the SiPMs boards, remaining almost constant at 17±1 °C. No significant variations in the photopeak channel were

SiPMs boards, remaining almost constant at 17 ± 1 °C. No significant variations in the photopeak channel were found due to temperature variations. The energy resolution calculated as $\Delta E(FWHM)/E_{centroid}$ was also monitored for regions of interest around the measured radioactive source (around 3 mm × 3 mm) and found to be stable with an average value of 13 ± 1 %, 14 ± 1 % and 17 ± 1 % for the 10, 15 and 20 mm thick blocks, respectively. An energy window of 15% (434-588 keV) was applied to all data.

3.1. Intrinsic resolution measurements

In the following we describe the two main set-up configurations used in this work, see Fig. 2 (a)-(b). The first set-up shows the method for which one detector is moved far away. The opposite detector is sequentially moved from the detector under study from a distance of 115 mm up to 1780 mm in order to measure the resolution as a function of the collimation angle. There is no significant influence of the source size on the determined

1 collimation angle (about 0.5% maximum) or due to the non-collinearity of the annihilation photons (maximum

2 0.3%) for the distances we have considered. In this case, the natural Line of Response (LOR) collimation angle is

3 a function of the distance between the two detectors and, therefore, can be calculated as:

$$\theta_{DD} = atan \frac{x}{s_d} \tag{1}$$

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where x is the half of the reference detector crystal size, 25 mm in our case, and S_d is the separation distance between the reference detector and the source. Here we did not perform any type of collimation. By moving away the reference detector we are "naturally" collimating the source, and therefore, isolating the source contribution to the measured spatial resolution FWHM of the detector under study. The acquisition time was increasing as a function of the separation distance between the modules in order to retrieve similar overall counts for all distances (~2×10⁵ coincidences). A coincidence window of 5 ns was selected. The data acquisition system has been described elsewhere, see for instance references [5][23]. The second set-up, as depicted in Fig. 2 (b), describes the method we are proposing. Here we fixed the distance between the two detector blocks to 115 mm. We acquired during 15 minutes recording ~2×10⁶ coincidences whereas in the previous method we need to acquire at least 90 minutes for the longest separation distance recording a factor ten less coincidences. In this case, software collimation is applied to the data. The software collimation varies as a function of the LOR angle, and consists on allowing only events whose LOR has a slope smaller than a predefined one. For each coincidence we calculate the angle between its LOR and the normal ($\theta_{software}$), see Fig. 2 (b). If $\theta_{software}$ is smaller than the selected collimation angle, the event is accepted. The system accepts events from squared regions under this angular restriction. Events in each detector block are binned in 600 × 600 pixels. In particular, we have configured collimation angles varying from 0.001 radians to nearly 0.2 radians from the normal, corresponding to squared regions in the reference detector of 0.23 mm and 46.60 mm, respectively. Smaller collimation angles were not considered because of the lack of statistics. In

contrast to the distant detectors set-up, in the proposed method only one measurement is needed.

To study the intrinsic detector resolution across the whole crystal surface, the source was displaced in steps of

1 mm from one border of the crystal to its center, assuming a symmetrical behavior of the other half. Each data

point was software collimated in the range of 0.001 rad to 0.2 rad.

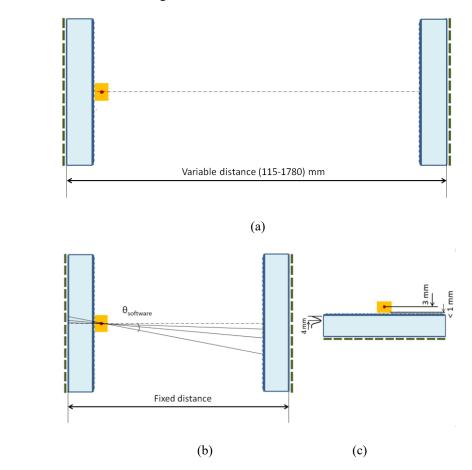


Fig. 2. Sketch of the experimental setups. Both detectors included a monolithic LYSO crystal with black painted lateral walls and were read out using an array of 16×16 SiPMs. (a) Distant detectors method showing the small size source of ²²Na in front of the detector under study. (b). Software Collimation method. (c) Sketch showing the distance from the source to the scintillator active volume. It is also depicted the most probable DOI, about 4 mm, in a 10 mm thick LYSO crystal.

3.2. Source simulation

Using GATE, a simulation platform based on the Geant4 Application for Tomographic Emission [34], we have carried out simulations of the ²²Na source profiles in order to deduce the expected contribution of the

positron annihilation range to the experimental measurements. The simulated source is a ²²Na sphere of 0.25 mm in diameter inside a PMMA disk (12.7 mm radius, 6 mm thick). In these simulations, in contrast to the default parameters in GATE for this source, the whole energy distribution of the ²²Na positron emission was considered and not only the maximum energy value. Figure 3 shows the energy spectra considered for the ²²Na source together with the spectra of other isotopes of interest in PET scanners. This distribution was calculated following an analytical expression valid for allowed or super allowed beta decays [27][35]. The theoretical energy distribution is of the form:

$$N(E)dE = gF(Z, E)pE(E_{max} - E)^{2}dE$$
 (2)

where N(E) is the number of decays at energy E, g is a coupling constant, E is the β particle energy, E_{max} is the maximum energy of that particles (0.545 MeV for 22 Na), p is its momentum, F(Z, E) is the Fermi function, and Z is the atomic number of the beta decay daughter. A non-relativistic approximation for the Fermi function, valid for allowed transitions of lighter elements is [38][39]:

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$$F_{allowed}(Z, E) = 2\pi \frac{\eta}{(1 - e^{-2\pi\eta})}$$
 (3)

where $\eta = -Z\alpha E/p$ for positron decays, and $\alpha = 1/137$, the fine structure constant.

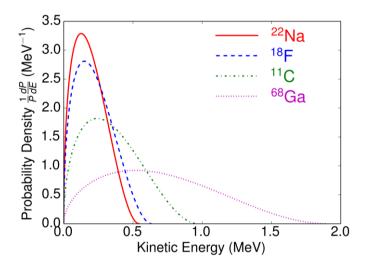


Fig. 3. β^+ -decay energy spectra for different isotopes of interest for PET studies. In particular, the solid red line shows the energy spectra of the 22 Na source used in this study. The energy curve for the 68 Ga has also been calculated (dotted pink line), as it is not also found by default.

The simulation contour plot and the profile of the annihilation points of the source are shown in Fig. 4 (a) and (b), respectively. We have performed fits to these profiles with both Gaussian and Lorentzian distributions.

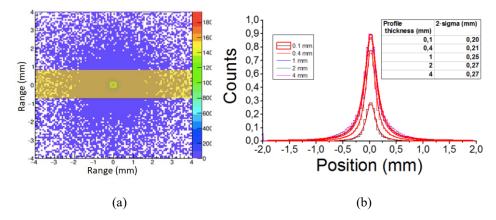


Fig. 4. (a) Contour plot for the simulated 22 Na 0.25 mm source. (b) Simulated profiles (see yellow band in panel (a)) and fits carried out using a Lorentzian distribution. Profiles for different band widths ranging from 0.1 mm to 4 mm were studied.

Better agreement is achieved when using Lorentzian functions, with 0.99 regression coefficients. The Lorentzian FWHM of the fitted profiles will be used in the following to estimate the intrinsic detector spatial resolution. The analysis of the experimental data is done by studying a 1D projection of the 2D projections of the annihilation point distribution. The thicknesses of the projection bands have been studied since those slightly impact the resulting Lorentzian FWHM, see Fig. 4. For band widths wider than 1 mm, the determined FWHM remains constant at 0.27 mm. For a very thin band with a width of only 100 µm, the difference from 4 mm bands is smaller than 30 %, see Table 1. We have finally selected a Lorentzian FWHM of 0.27 mm for the ²²Na source, which corresponds to the value found for profile thicknesses 2-4 mm.

Table 1. Lorentzian FWHM of the source profile for different band widths.

Band width (mm)	FWHM (mm)	
0.1	0.20	
0.4	0.21	
1	0.25	
2	0.27	
4	0.27	

3.3. Data analysis

 During the data analysis we have characterized the measured profiles of the small size source. In order to provide accurate determination of the width of the measured profiles, those need to be given in metric units. For this purpose, the radioactive source was displaced across the crystal surface at known positions. The observed source in the flood maps are projected onto the X and Y axis, see example in Fig. 5 (a)-(b). We have projected bands of several pixels, typically 10 pixels corresponding to approximately 1.6 mm, onto the X and Y axes. Smaller bands widths have also been tried with insignificant differences in the results.

The relation between the measured position in channels and the known geometrical beam position fits well to a first order polynomial, in the central crystal region. However, in order to account for the truncation of the light distribution at the crystal edges that renders compression of the events in this region, a third order polynomial fit is better suited [23]. Figure 6 (a), (b) and (c), show the mechanical curve versus the measured positions obtained for the three different scintillators. The thicker crystal (20 mm) suffers from a larger truncation of the data at the crystal edges, which explains why the 20 mm thick crystal curve shows a more accused compression. The plots show the bias, which is the deviation between the mechanical and the measured source position, across the crystal surface. For the 10 mm thick crystal, the bias is smaller than in the 15 mm and 20 mm thick crystals, as expected. For the thickest crystal case, due to the stronger light truncation, the bias is larger in the extrema, about 6 mm. However, these effects do not constrain the current results allowing characterizing events in the whole surface.

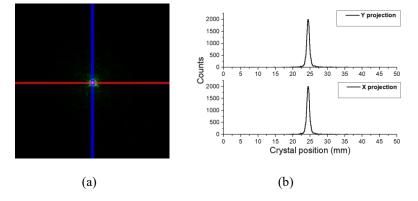


Fig. 5. (a) Flood map of the image with the point-like source in the center of the 10 mm thick block 600×600 pixels. (b) Two projections (X and Y) for the blue and red bands depicted in (a).

1 BIAS 10mm 8 7 BIAS (mm) 2 --2 -6] -8 1 -25 20 -15 -10 -5 0 5 10 15 20 Mechanical source position (mm) 25 2 3 (a) BIAS 15mm 8 -6 4 -BIAS (mm) 2 0 --2 -6 -8 20 -15 -10 -5 0 5 10 15 20 Mechanical source position (mm) 4 5 (b) BIAS 20mm 8 6 4 BIAS (mm) 2 0 -2 -6 -8 -20 -15 -10 -5 0 5 10 15 20 Mechanical source position (mm) 6 7 (c)

Fig. 6. Deviation between the known mechanical and measured source position across the crystal surface for (a) the 10 mm, (b) the 15 mm, and (c) the 20 mm thick blocks, respectively.

3.4. Extracting the intrinsic resolution

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The main objective is to provide a model that can be used to determine the detector intrinsic resolution by fitting the measured source profiles. The detector intrinsic spatial resolution is convolved with the source dimension resulting on the measured $FWHM_{measured}$. If both, detector and source, were to follow Gaussian distributions, the resulting $FWHM_{measured}$ could be expressed as $\sqrt{FWHM_{source}^2 + FWHM_{detector}^2}$, which is the usual way of dealing with this entangle. Indeed, this approach works well when the source dimensions are large and therefore closer to a Gaussian distribution. However, in this work we aim to isolate as much as possible the source contribution from the detector under analysis and, therefore, a very small size source was used (0.25 mm in diameter). Assuming that the positron-emitter source follows a Lorentzian profile and the detector intrinsic resolution follows a Gaussian distribution, it turns out that the convolution of these distributions is the so-called Voigt function [36], which cannot be evaluated in closed form and therefore has to be evaluated numerically. However, in our case there is a collimation effect that produces a broadening of the source distribution. For that reason the convolution of the detector Gaussian shaped distribution and the "collimated" source profile is no longer a Voigt distribution. Therefore, the profiles of the projected 1D collimated beam images (see Fig. 5b) contain three essential contributions namely, i) the distribution of the gamma emission points in the positron source; ii) the influence of the gamma beam created by the software collimation; and iii) the intrinsic detector resolution. To disentangle the effect of these three components, a mathematical model of the measured beam profiles was developed which takes into account the contribution of each of them. In this work we assume that the intrinsic resolution PDF is Gaussian with a FWHM characterizing the intrinsic resolution. This intrinsic resolution PDF has to be convolved with the normalized intensity distribution of the gamma emission points in the ²²Na point source and the software collimation method.

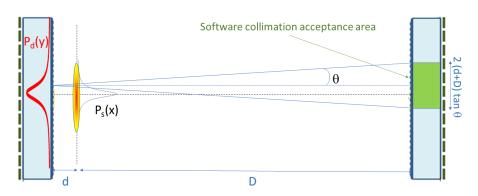


Fig. 7. Schematic drawing showing the nomenclature employed in the mathematical beam profile model used to extract the intrinsic resolution.

3.4.1 Combined influence of source distribution and software collimation

The probability that a gamma emitted by the emission point distribution $P_s(x)$ of the ²²Na source hits the detector surface at a point y at a distance d from the source (see Fig. 7), while fulfilling the software collimation constraint (i.e. the angle between the LOR connecting the impact point on the detector under study and the 2nd detector at a distance D from the source used to perform the software collimation, is smaller than a given collimation angle θ) can then be expressed as:

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$$P_d(y) = \frac{1}{C} \int_{-\infty}^{\infty} P_s(x) \cdot B\left(y + (x - y) * \frac{d + D}{d}, y, 2 * (d + D) * \tan(\theta)\right) dx$$
 (4)

where θ is the maximum photon incidence collimation angle allowed in the monolithic block, C is a constant defined such that $P_d(y)$ is a PDF and therefore its integral over all possible y values should be 1 and the function B(x, c, w) represents the software collimation effect and is defined as an unit box function with its center at c and has a width w, that can be defined as:

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$$B(x,c,w) = \begin{cases} 0 & \text{if } x < c - w/2 \\ 1 & \text{if } c - w/2 \le x \le c + w/2 \\ 0 & \text{if } c + w/2 < x \end{cases}$$
 (5)

In section 3.2 it has been shown that the 1D gamma emission point distribution $P_s(x)$ of the ²²Na source used in the experimental set-up can be modelled with a Lorentzian distribution with a FWHM_{source} of 0.27 mm.

$$P_S(x) = \frac{2}{\pi \cdot FWHM_{source} \left(1 + \frac{4x^2}{FWHM_{source}^2}\right)}$$
 (6)

- 1 Evaluating the integral in eq. 4 then can be shown that the effect of the software collimation procedure
- 2 produces a 1D Probability Density Function (PDF) of photons impact position y on a plane at a distance d from
- 3 the source that can be expressed as:
- 4 $PDF(y|d, FWHM_{source}, \theta) =$

$$5 \quad \frac{1}{\pi \cdot K} \cdot \left(tan^{-1} \left(\frac{2y}{FWHM_{source}} + \frac{2d \tan(\theta)}{FWHM_{source}} \right) - tan^{-1} \left(\frac{2y}{FWHM_{source}} - \frac{2d \tan(\theta)}{FWHM_{source}} \right) \right) \quad (7)$$

6 where K is a normalization constant given by:

$$7 K = \int_{-\infty}^{\infty} \frac{1}{\pi} \left(tan^{-1} \left(\frac{2(y + d \tan(\theta))}{FWHM_{source}} \right) - tan^{-1} \left(\frac{2(y - d \tan(\theta))}{FWHM_{source}} \right) \right) = 2 d tan(\theta)$$
 (8)

- We will refer to θ as $\theta_{software}$ when referring to the software collimation method and θ_{DD} for the distant
- 9 detector method. Since the gammas do not interact at the surface of the detector but penetrate to different depths,
- 10 the final gamma interaction distribution in the detector is given by a weighted average of the
- 11 $PDF(y|d, FWHM_{source}, \theta)$ at different interaction depths z in the detector:

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$$\int_{d_{min}}^{d_{max}} e^{-\mu \cdot (z - d_{min})} * PDF(y|d, FWHM_{source}, \theta) \cdot dz \qquad (9)$$

- In the above equation, it was assumed that the front of the detector is at a distance of d_{min} from the source
- while d_{max} is equal to d_{min} + detector thickness. Therefore, the d parameter takes into account the distance from
- 15 the source to the impact position within the scintillator material. The average interaction depth \bar{d} within the
- 16 crystal can be theoretically estimated as a cone with opening angle θ [37]:

$$\bar{d} = \frac{\cos\theta}{\mu} + \frac{L}{1 - e^{\frac{\mu L}{\cos\theta}}} \tag{10}$$

- where μ is the attenuation coefficient of the LYSO material at 511 keV (0.87 cm⁻¹), and L is the crystal thickness.
- We obtain values of \bar{d} of about 4.2 mm, 5.6 mm and 7.2 mm from the entrance surface for the 10 mm, 15 mm
- and 20 mm thick crystals, respectively. Since there are 3 mm source encapsulation (half thickness) + 0.5 mm
- housing + 0.5 mm retroreflector $+ \sim 0.5$ mm spacing between the source edge and the detector housing, see Fig. 2
- 22 (c), d is estimated to be about 9 ± 1 mm, 10 ± 1 mm and 12 ± 1 mm for each crystal respectively.

3.4.2 Software collimation model limitations

In the model, to predict the gamma interaction point distribution probability at a given distance d from the point source, the effect of multiple interactions (i.e. a Compton scatter followed by a photoelectric absorption or a second Compton scatter in the same block) is neglected. However, in a real experiment, gammas that undergo multiple interactions will have a measured position that is usually (much) further away from the collimated beam center in comparison to single interaction events. When small software collimation angles are used, many of these multiple interaction events will be rejected for analysis and hence the model remains valid for most events in a measured data set.

Regarding gamma-rays impinging closer to the crystal edges, i.e. when the collimated beam is moved very close to the edge of the two opposite blocks, the gamma interaction point distribution probability gets truncated at one side. This effect was not put in the current model because no analytical solution of equation 7 could be found in this case. Hence the contribution of the beam collimation in the last 1 or 2 mm from the edge can be slightly overestimated. However, for small collimation values there is almost no influence of this effect on the data and therefore, since the model takes into account all the collimated profiles we do not expect a significant impact in the final resolution value.

3.4.3 Adding the intrinsic resolution contribution to the measurement model

To model the final profile of the measured data we convolved the source Lorentzian modified profile (i.e. $PDF(y|d, FWHM_{source}, \theta)$) with the Gaussian intrinsic detector resolution point spread function (PSF), characterized by a standard deviation σ . This resulted on the final resolution model given by:

20
$$f(xy|d, FWHM_{source}, \theta, \sigma) =$$

$$21 \qquad \frac{\cot\theta}{2\pi d} \cdot \sum_{n=0}^{1} (1-2n) \cdot \left(\frac{\pi}{2} \operatorname{Re}(\operatorname{Erf}(z_n)) - \operatorname{Im}\left\{z_{n-2}^2 F_2(1,1; \frac{3}{2}, 2; -z_n^2\right\}\right) \tag{11}$$

22 with

$$z_n = \frac{2d \cdot (1-2n) \cdot \tan(\theta) + i \cdot FWHM_{source} + 2y}{2\sqrt{2}\sigma}$$
 (12)

Re and Im are operators that take the real and imaginary parts respectively. Erf and ${}_2F_2$ represent the Error and a generalized hypergeometric functions, respectively. This equation was fitted simultaneously to measure the resolution profiles taken at eight different collimation angles (from 0.009 up to 0.08 radians) where the effect of the different depth of interactions on the final distribution can be considered negligible. For large collimation angles, the shape of the weighted average PDF is very different compared to the PDFs at a specific distance. Hence, the deduced equation cannot be used reliably in these cases. For small to medium collimation angles the shape of the weighted average PDF remains similar to the shape of the PDFs at a specific distance. Since the intrinsic spatial resolution is independent of the collimation angle, σ was taken as a common parameter in the simultaneous fits. $FWHM_{source}$ was fixed at 0.27 mm, as was derived from simulations, while the distance d was left as a free parameter to fit in order to verify if its value would converge close to the expected average interaction depths.

4. Results

4.1. Validation of the software collimation method

We have first validated the proposed method of software collimating the data with the most traditional one of sequentially moving backwards the detector for coincidences. The validation test was carried out for the 10 mm thick LYSO crystal and with the source positioned at its center.

Figure 7 shows the $FWHM_{measured}$ as a function of the collimation angle for both the distant detector approach (open circles), and the proposed software collimation method (full squares). Notice that we have not carried out any photon DOI selection to the data. We selected projections in the flood maps (600×600 pixels) of the source onto the X and Y axes. Each data point is calculated as the average of the $FWHM_{measured}$ for the X and Y profiles. The $FWHM_{measured}$ is obtained by fitting the measured data to Gaussian distributions. This is a good approximation to the alternative way of calculating the maximum value of the distribution and measuring the width at the half of that maximum for each profile. We have checked this assumption by manually measuring the

width of several profiles. Good agreement was obtained for all crystal geometries, obtaining deviations smaller than 2% and, thus, validating the Gaussian fitting procedure. The error bars associated to each point are calculated as the standard deviation of the two (X and Y) Gaussian FWHM to each profile.

A good agreement between the two collimation methods was observed, as depicted in Fig. 8. In the following section both data sets have been fitted using the proposed model, retrieving identical detector intrinsic resolution values. Notice that whereas in the distant detectors method the number of counts is similar for all collimation angles, in the software collimation method the statistics are being reduced as a function of the angle due to the reduction in the accepted number of counts.

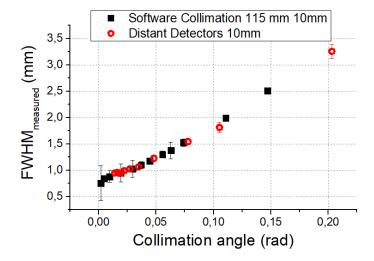


Fig. 8. Validation of the proposed method (software collimation, full squares) compared to the distant detector method (open circles) for the 10 mm thick crystal.

4.2. Gamma interaction distribution and source modelling

As we have already pointed out, the gammas do not interact at the detector surface but penetrate to different depths within the scintillator block. Therefore, the final gamma interaction distribution in the detector is modelled using Eq. 9. Figure 9 shows the modelled source distribution in the detector for the 10 mm thick block. These results show the PDFs for different collimation angles and at different depths overlapping with the weighted average PDF (black solid line, assuming the front of the detector is 4.5 mm away from the source, i.e. d_{min} =4.5 mm in Eq. 9). For large collimation angles, the shape of the weighted average PDF is very different compared to

the PDFs at a specific distance. Hence, Eq. 9 cannot be used reliably in these cases. However, for small and medium collimation angles the shape of the weighted average PDF remains similar to the shape of the PDFs at a specific distance. For this reason we are considering angles ≤ 0.1 radians for the model.

To find out the average interaction distance at which Eq. 9 matches best with the weighted averaged PDF, we have plotted the variation of the integrated absolute difference between the weighted averaged PDF and PDFs at a specific depth. Figure 10 shows these values for two different collimation angles. For the 10 mm thick crystal the best agreement is achieved at a depth of 9±1 mm from the source, corresponding to approximately 5±1 mm DOI from the crystal entrance. Using Eq. 10 we estimated an average interaction distance of 9±1 mm (i.e. 4.2 mm DOI), which agrees well with the obtained value.

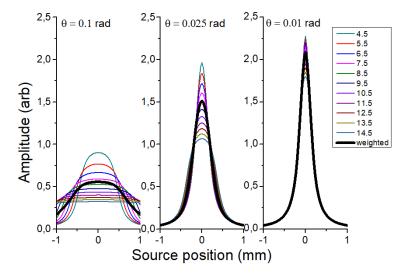


Fig. 9. PDFs at individual depths (in mm) in comparison with the weighted average PDF (black solid line) for three different collimation angles (in radians). The profiles are obtained for the 10 mm thick block.

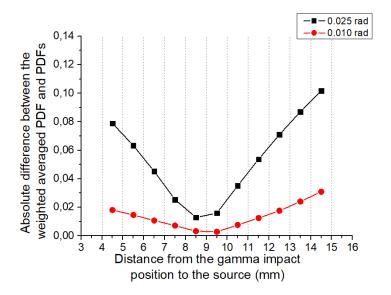


Fig. 10. Integrated absolute difference between the averaged PDF and a PDF at a specific interaction distance from the source for the 10 mm thick block.

4.3. Modelling of the measured resolution profiles. Fitting analysis

We have fitted the measured source profiles for the different collimation angles using Eq. 11 and Eq. 12. The fitting returns the source-detector distance, the $FWHM_{source}$ and the detector intrinsic resolution. Notice that Eq. 11 is not completely valid for large collimation angles, so we avoid using profiles whose collimation angle is \geq 0.1 radians. The profiles for all collimation angles are simultaneously fitted using Eq. 11, sharing the free detector intrinsic resolution parameter. The source size contribution ($FWHM_{source}$) was also shared but fixed at 0.27 mm, as earlier predicted by simulations. The data obtained for the smallest collimation angle was not included due to the lack of statistics. The number of angular samples considered for the fitting was 6; using \pm 1 samples we did not observe significant changes. Table 2, summarizes the angular range considered for each crystal.

Table 2. Angular range considered for each studied crystal.

Scintillator thickness (mm)	Angular range (radians)	
10	0.009-0.08	
15	0.009-0.08	
20	0.009-0.07	

We have first validated the model with the measurements performed at the crystals center. The fitting of both the X and Y profiles was performed using Wolfram Mathematica [40]. The fitting procedure yields a detector intrinsic resolution FWHM of 0.64 ± 0.02 mm for the 10 mm thick crystal when using the software collimation data. The data obtained when using the distant detector set-up was also analyzed using the same methodology resulting on a comparable detector intrinsic resolution FWHM of 0.65 ± 0.03 mm, therefore, confirming the robustness of the software collimation set-up. Once the software collimation method was validated with the 10 mm thick block, identical data sets were acquired using the software collimation set-up for the 15 mm and 20 mm thick blocks. The analysis of the data returned values of 0.82 ± 0.02 mm and 1.07 ± 0.03 mm FWHM, respectively. Note that the detector intrinsic resolution FWHM was estimated as the average value of both the X and Y profiles. The quoted uncertainty corresponds to the standard deviation between the X and Y profiles.

These analyses also provided information on the distance between the source and the interaction point of 7.9 ± 0.2 mm, 9.0 ± 0.3 mm and 11.1 ± 0.4 mm for the 10 mm, 15 mm, and 20 mm thick blocks, respectively. These results agree well with the expected values estimated using Eq. 10. Figure 11 shows in black circles the measured profiles and with a red line the model prediction for the 10 (top), 15 (central) and 20 (bottom) mm thick blocks, and for three different collimation angles. A good agreement between the measured profiles and the model is observed for each collimation angle.

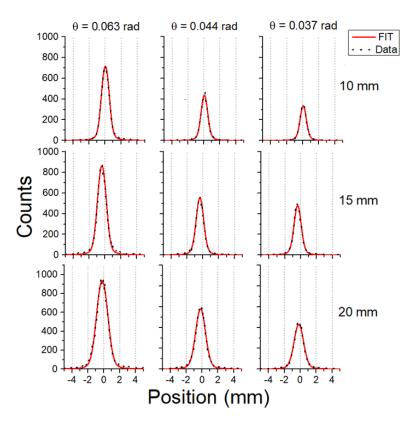


Fig. 11. Measured profiles (black points) and model prediction (red line) for three collimation angles (in radians). Top, profiles for the 10 mm thick crystal. Central. Profiles for the 15 mm thick crystals. Bottom, profiles for the 20 mm thick crystal.

4.4. Impact position dependence

The acquired profiles have been analyzed using the proposed model in order to obtain the dependence of the detector intrinsic spatial resolution with the impact position. Here, the FWHM_{source} was again fixed to 0.27 mm and d was left as a free parameter. Figure 12 shows the detector intrinsic spatial resolution FWHM as a function of the mechanical source position. In order to take into account the influence of the crystal edges and since we moved the source across the x-axis of the detector, we have calculated the resolution values for the X profiles of the source. The variation of the calculated detector intrinsic spatial resolution FWHM with the impact position is larger for the 15 and 20 mm thick crystal cases. This is produced because the thicker crystals produce a stronger truncation of the LD and, thus, a worse characterization of its centroid position (see Fig. 6) [11].

Table 3. Intrinsic detector spatial resolution (FWHM_{detector}) obtained for all scintillator geometries.

Scintillator	Central	Average	Average
	Source	[-20,20] mm	[-25,25] mm
10 mm	0.64 ± 0.02	0.67±0.03	0.68±0.04
15 mm	0.82 ± 0.02	0.85 ± 0.06	0.90 ± 0.14
20 mm	1.07 ± 0.03	1.22 ± 0.10	1.29±0.19

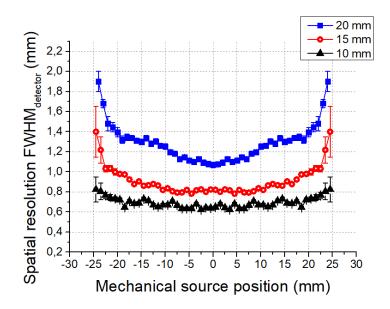


Fig. 12. Detector intrinsic spatial resolution FWHM as a function of the source mechanical position. Black full triangles show the results for the 10 mm thick crystal, the red open circles those for the 15 mm thickness and blue full squares are for the 20 mm thick block.

For data corresponding to the whole crystal surface, the mean detector intrinsic spatial resolution was found to be 0.68 ± 0.04 mm for the 10 mm case, slightly improving to 0.67 ± 0.03 mm when the range is reduce to [-20, 20] mm, since less truncation is observed. The 15 mm thick crystal exhibits an average intrinsic spatial resolution for the whole crystal of 0.90 ± 0.14 mm. This improves to 0.85 ± 0.06 mm when the range is reduce to [-20, 20] mm. Finally, the 20 mm thick crystal exhibits an average intrinsic spatial resolution for the whole crystal of 1.29 ± 0.19 mm. This improves to 1.22 ± 0.10 mm when the range is reduced to [-20, 20] mm. Table 3 summarizes the average values for impacts only at the crystal center, for the range varying from -20 to 20 mm and when considering the whole crystal surface from -25 to 25 mm.

5. Discussion and conclusions

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We have proposed a method to determine the intrinsic detector spatial resolution in detector blocks based on monolithic crystals suitable for PET systems. This approach can be done while the PET system is already mounted and only requires one measurement. The method consists on software collimation of the measured data in order to isolate the finite source size FWHM_{source} contribution to the FWHM_{measured}. We have derived Eq. 11 that uses this information and returns the intrinsic detector spatial resolution together with the most suitable distance between the source and the interaction depth within the scintillation block. The method has been validated against the approach of using a bench-top set-up where the reference detector is sequentially moved backwards, requiring a higher number of measurements and also a long separation distance between the detectors in order to avoid the influence of the source size on the reference detector. We have found comparable results when using the two experimental approaches, i.e. the proposed software collimation method and the geometrical collimation using a bench set-up ("distant detector" approach), see Fig. 7. This new experimental method allows obtaining the detector intrinsic spatial resolution by performing only one measurement. However, this measurement should have enough statistics to allow applying small angular restrictions without introducing significant statistical fluctuations. The proposed method is capable to provide the intrinsic detector spatial resolution within an assembled PET scanner. In this work results for a fixed distance of 115 mm are shown regarding the case of the software collimation method. However, other distances have also successfully been tested for this method retrieving the same results. Averaged intrinsic detector spatial resolutions for the whole crystal surface, of 0.68±0.04 mm, 0.90±0.14 mm and 1.29±0.19 mm have been obtained for the 10 mm, 15 mm and 20 mm thick crystals, respectively. The intrinsic spatial resolution strongly depends on the detector block characteristics, such as the scintillator geometry and the photosensor sampling matrix. In this case we are using a 16×16 SiPM matrix, 3.26 mm pitch. For SiPM arrays with larger pitches or larger photosensor sizes we expect some degradation on the intrinsic spatial resolution. As shown along this work, also the crystal geometry has an impact on the $FWHM_{measured}$, the thicker the crystal the worse the $FWHM_{measured}$.

The measured profiles were fitted using a simplistic version of the model based on the so-called Voigt function (see section 3.4). The main drawback of this model is that did not account for the influence of the software collimation in the profiles and required *d* to be fixed for events towards the edges of the detector. An underestimation of the detector intrinsic resolution of about 10% was observed when compared with the proposed method [41]. Other works are based on determining the point spread function (PSF) of the whole scanner system at the imaging space, (mainly based on Gaussian fits [42]) or at the projection space. In the last case, the system response function can be determined through analytical derivations, Monte Carlo simulations, or empirically [43]. In theory, the empirical approach of physically measuring the system response to a collimated point source leads to the most accurate description of the system [44]. However, as far as we know, no empirical determination of the intrinsic spatial resolution at individual detector module elements has been reported. The accurate knowledge of this individual intrinsic spatial resolution and its dependency with the impact position in the monolithic detector surface allows one considering this information during the reconstruction process. For instance, one could include this information in the reconstruction algorithms, expecting to produce an improvement on the quality of the final image.

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