



Comprehensive Analysis of Alpha-Parametric Set for the Calculation of Intersection Lengths of Radiological Ray Path in Siddon's Algorithm Used in 3D Image Reconstruction

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Abstract – The Siddon algorithm is one of the radiological ray path (x-ray) calculation tools used in 3D image reconstruction in medical imaging. In the algorithm, a set of alpha-parametric values is computed containing the length and index values where the voxel array of the x-ray intersects the x-y-z axes. In the alpha-set creation section of the Siddon algorithm, the set elements are sorted from small to large, but some elements have been noticed to have the same value in simulations. These elements are used to calculate which voxels are hit by the ray along the radiological path and at what ratio, but it was recognized that some values of the set were zero, which means some rays did not intersect some voxels at all. This situation may lead to data loss in 3D image reconstructions in medical imaging such as digital breast tomosynthesis (DBT) and computed tomography (CT) especially for huge dimensions such as size up to 800×800×50. Considering the mentioned problems, in this study, the effect of using or eliminating the same repetitive values in the alpha parametric set of the Siddon algorithm on calculations was investigated. To prove our proposal, we performed 3D image reconstruction (lossless and lossy) of a synthetic phantom at a size of 100×100×50. Using special functions that do not take into account the duplicate values in the algorithm, excluding the duplicate values from the calculation solved the stated problems (lossless reconstruction). In this way, data loss that may occur in 3D image reconstruction was reduced since voxel indices and intersection lengths were matched correctly and meaningfully.

Keywords – Computed tomography, digital breast tomosynthesis, siddon algorithm, x-ray, 3D image reconstruction

1. Introduction

Mainly, iterative and analytical image reconstruction methods are applied in medical imaging modalities such as the optoacoustic tomography (OAT) (or photoacoustic tomography (PAT)) (Paltauf et al., 2002; Wang et al., 2012), the optical tomography (OT) (Dekker, Battista, & Jordan, 2017; Klose & Hielscher, 1999), the diffuse optical tomography (DOT) (Üncü et al., 2017; Sevim et al., 2017; Mercan et al., 2017; Mercan et al., 2019), the computed tomography (CT) (Sidky et al., 2006; Biguri et al., 2017) and the digital breast tomosynthesis (DBT) (Oliveira et al., 2016; Polat & Yildirim, 2018; Polat et al., 2019a). The CT, developed by Hounsfield and Cormack in the 1970s (Nobel Media AB, 2014; Raju, 1999), enables three-dimensional (3D) images of the examined part of the body like the head, chest, and breast. The DBT, an advanced form of x-ray mammography evolved by Gershon-Cohen and others (Kopans, Meyer, & Sadowsky, 1984), focuses on the examination of the breast by scanning at a limited angle (Helvie, 2010; Niklason et al., 1997; Wu et al., 2004). For reconstructing three-dimensional (3D) images of the part of the body, the algebraic reconstruction technique (ART) introduced by (Kaczmarz, 1937) and the simultaneous ART (SART) introduced by (Andersen & Kak, 1984) have been used as iterative algorithms, while the filtered back projection (FBP) which was first advanced by Riddle and Bracewell in 1967 (Bracewell & Riddle, 1967) and independently in 1971, by Lakshminarayanan and Ramachandran (Ramachandran & Lakshminarayanan, 1971) has been used as an analytical algorithm, (Kak, Slaney, & Wang, 2002). Whether it is an iterative or analytical method, in

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modalities such as OAT, OT, DOT, CT and DBT, the geometry of the x-rays sent to the target object should be calculated for 3D image reconstruction system design.

The Siddon algorithm (Siddon, 1985), which effectively calculates the ray path geometry, can be used to calculate the system matrix elements in constructing the 3D image reconstruction space. The Siddon algorithm was applied in many applications in ray-path calculation studies and in medical imaging (Gao, 2012; Jacobs et al., 1998; Li et al., 2008; Polat & Yildirim, 2018; Polat et al., 2019a; Xue, Zhang, & Pan, 2011; Zhao & Reader, 2003). There are six main sections in the Siddon algorithm including calculating the range of alpha-parametric values, calculating the range of the indices at x-y-z axes, calculating the alpha-parametric sets for x-y-z axes, merging sets to form alpha-set, calculating voxel intersection length, and calculating voxel indices, respectively.

In the alpha-set creation section of the Siddon algorithm, the alpha-set elements are sorted from small to large, but some elements have the same value or zero-value. The values in the set are used to calculate the voxel intersection length and voxel indices, therefore it is a critical issue to consider whether eliminating some values of zero and/or identical or not. This situation may lead to data loss in the applications of the Siddon algorithm for 3D image reconstructions in medical imaging such as DBT and CT. Considering the mentioned problems, in this study, we analyzed the creation steps of the alpha-parametric set that affect the calculations of voxel intersection length and voxel indices and investigated the effect of using or eliminating the repetitive and/or zero- values for the calculations in the Siddon algorithm.

2. Materials and Methods

The Siddon algorithm introduced an efficient, reliable, and exact calculation of a radiological route through a 3D-array of CT (Siddon, 1985). The schematic of two-dimensional (2D) representation for the 3D CT array of (N_x, N_y, N_z) voxels is given in Figure 1. A single ray starting from point $P_1(x_1, y_1)$ to point $P_2(x_2, y_2)$ passes through the CT array. The radiological path in 3D space, in other words, the ray sum is defined by 2.1.

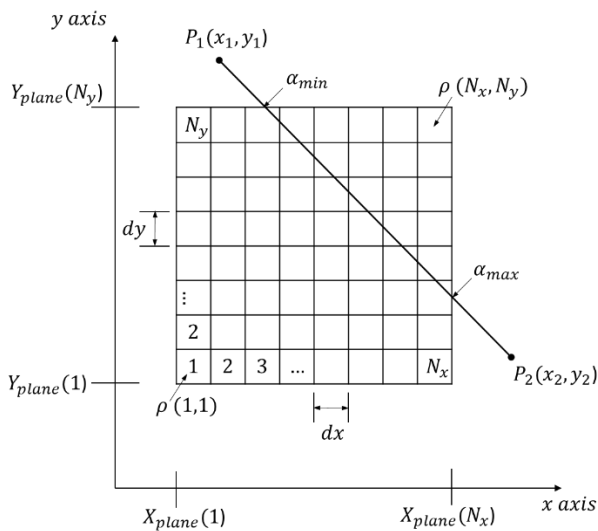


Figure 1. The schematic of two-dimensional (2D) representation for the 3D CT array of (N_x, N_y, N_z) voxels

$$d = \sum_i \sum_j \sum_k l(i, j, k) \rho(i, j, k) \tag{2.1}$$

where $l(i, j, k)$ is the voxel intersection length and $\rho(i, j, k)$ is the density of voxel in the point (i, j, k) .

In the Siddon algorithm, the ray from $P_1(x_1, y_1, z_1)$ to $P_2(x_2, y_2, z_2)$ is described with the alpha-parametric set; α ($\alpha_1 = 0$ at P_1 and $\alpha_2 = 1$ at P_2) in 3D space (see 2.2).

$$\begin{aligned}
 X(\alpha) &= X_1 + \alpha(X_2 - X_1) \\
 Y(\alpha) &= Y_1 + \alpha(Y_2 - Y_1) \\
 Z(\alpha) &= Z_1 + \alpha(Z_2 - Z_1)
 \end{aligned}
 \tag{2.2}$$

Alpha-parameter (α) set in the range of $(\alpha_{min}, \alpha_{max})$ indicates the intersections of the ray with planes. If P_1 or P_2 are within the grid boundaries, then α_{min} gets zero, and α_{max} gets one, otherwise they become a nonzero value (see Figure 1).

For the (N_x, N_y, N_z) voxel array, orthogonal sets of equidistant parallel planes are written as in 2.3.

$$\begin{aligned}
 X_{plane}(i) &= X_{plane}(1) + (i - 1)dx & i &= 1, 2, \dots, N_x \\
 Y_{plane}(j) &= Y_{plane}(1) + (j - 1)dy & j &= 1, 2, \dots, N_y \\
 Z_{plane}(k) &= Z_{plane}(1) + (k - 1)dz & k &= 1, 2, \dots, N_z
 \end{aligned}
 \tag{2.3}$$

where $dx, dy,$ and dz are the dimensions of the voxel and the distances between the x, y, z planes.

The α -parametric values are calculated by 2.4 using 2.2 and 2.3.

$$\begin{aligned}
 & \text{if } X_2 \neq X_1, \\
 \alpha_x(1) &= [X_{plane}(1) - X_1]/(X_2 - X_1) \\
 \alpha_x(N_x) &= [X_{plane}(N_x) - X_1]/(X_2 - X_1)
 \end{aligned}
 \tag{2.4}$$

Expressions are similar for y and z planes for $\alpha_y(1), \alpha_y(N_y), \alpha_z(1),$ and $\alpha_z(N_z)$. After that α_{min} and α_{max} are calculated by 2.5.

$$\begin{aligned}
 \alpha_{min} &= \max\{1, \min[\alpha_x(1), \alpha_x(N_x)], \min[\alpha_y(1), \alpha_y(N_y)], \min[\alpha_z(1), \alpha_z(N_z)]\} \\
 \alpha_{max} &= \min\{1, \max[\alpha_x(1), \alpha_x(N_x)], \max[\alpha_y(1), \alpha_y(N_y)], \max[\alpha_z(1), \alpha_z(N_z)]\}
 \end{aligned}
 \tag{2.5}$$

In the case α_{min} is greater than or equal to α_{max} , the voxels are not intersected by the ray. α_{min} and α_{max} values are used to calculate the indices range of $(i_{min}, i_{max}), (j_{min}, j_{max})$ and (k_{min}, k_{max}) , which intersect the planes (2.6).

$$\begin{aligned}
 & \text{if } (X_2 - X_1) \geq 0, \\
 i_{min} &= N_x - [X_{plane}(N_x) - \alpha_{min}(X_2 - X_1) - X_1]/dx \\
 i_{max} &= 1 + [X_1 + \alpha_{max}(X_2 - X_1) - X_{plane}(1)]/dx \\
 & \text{if } (X_2 - X_1) \leq 0, \\
 i_{min} &= N_x - [X_{plane}(N_x) - \alpha_{max}(X_2 - X_1) - X_1]/dx \\
 i_{max} &= 1 + [X_1 + \alpha_{min}(X_2 - X_1) - X_{plane}(1)]/dx
 \end{aligned}
 \tag{2.6}$$

The similar notations are written for $j_{min}, j_{max}, k_{min},$ and k_{max} . Using these indices, the sets of alpha values $(\{\alpha_x\}, \{\alpha_y\}, \{\alpha_z\})$ which represent the intersections of the ray with the x, y, z planes are calculated as in 2.7 for $\{\alpha_x\}$, and with similar expressions for $\{\alpha_y\}$, and $\{\alpha_z\}$.

$$\begin{aligned}
 & \text{if } (X_2 - X_1) > 0, \\
 \{\alpha_x\} &= \{\alpha_x(i_{min}), \dots, \alpha_x(i_{max})\} \\
 & \text{if } (X_2 - X_1) < 0, \\
 \{\alpha_x\} &= \{\alpha_x(i_{max}), \dots, \alpha_x(i_{min})\}
 \end{aligned}
 \tag{2.7}$$

The definite intersections are determined by merging the sets $(\alpha_{min}, \alpha_{max}), \{\alpha_x\}, \{\alpha_y\},$ and $\{\alpha_z\}$ as alpha-parametric set. The merged alpha-parametric set $\{\alpha\}$ that has $n + 1$ elements is expressed as in 2.8.

$$\begin{aligned} \{\alpha\} &= \{\alpha_{min}, merge[\{\alpha_x\}, \{\alpha_y\}, \{\alpha_z\}], \alpha_{max}\} \\ \{\alpha\} &= \{\alpha(0), \alpha(1), \dots, \alpha(n)\} \end{aligned} \quad (2.8)$$

The difference of two consecutive terms in the alpha set is used to calculate the intersection length $l(m)$ in 2.9. d_{12} in 2.9 is the Euclidian distance (the total length) of the ray from P_1 to P_2 , and the formulation of it is given in 2.10. Once the $\{\alpha\}$ array is formed, the exploring of the voxels that are hit with the ray and the calculations of their intersection lengths are possible. The m th and $(m - 1)$ th values of the $\{\alpha\}$ array, which are the parametric weights of the intersected voxels, form the intersection length as in 2.9.

$$l(m) = d_{12}[\alpha(m) - \alpha(m - 1)] \quad \text{for } m = 1, 2, 3, \dots, n \quad (2.9)$$

$$d_{12} = \sqrt{(X_2 - X_1)^2 + (Y_2 - Y_1)^2 + (Z_2 - Z_1)^2} \quad (2.10)$$

The voxel indices $[i(m), j(m), k(m)]$ which intersected by the ray are calculated using the midpoint of two consecutive intersections (m th and $(m - 1)$ th alpha) as in 2.11.

$$\begin{aligned} i(m) &= 1 + [X_1 + \alpha_{mid}(X_2 - X_1) - X_{plane}(1)]/dx \\ j(m) &= 1 + [Y_1 + \alpha_{mid}(Y_2 - Y_1) - Y_{plane}(1)]/dy \\ k(m) &= 1 + [Z_1 + \alpha_{mid}(Z_2 - Z_1) - Z_{plane}(1)]/dz \end{aligned} \quad (2.11)$$

where the α_{mid} is the midpoint of the m th and $(m - 1)$ th alpha and is given by 2.12.

$$\alpha_{mid} = [\alpha(m) + \alpha(m - 1)]/2 \quad (2.12)$$

Finally, the projection value, P_θ , which is the radiological route can be obtained by summing over the n elements of the length array $l(m)$ obtained from $(n + 1)$ elements of $\{\alpha\}$ -parametric set multiplied by the densities of the voxels as in 2.13.

$$P_\theta = \sum_{m=1}^n l(m)\rho[i(m), j(m), k(m)] \quad (2.13)$$

where $\rho[i(m), j(m), k(m)]$ is the m th voxel density and θ is the projection angle.

The intersection lengths $l(m)$ for each ray are used to build up a system matrix \mathbf{A} for all projections (in this study, three projections, $\theta = -25^\circ, 0^\circ, +25^\circ$), and this system matrix \mathbf{A} is used to provide a solution for linear algebraic equation system given by 2.14, which models 3D image reconstruction. In 2.14, P_θ is the projections acquired from a target via a medical imaging modality such as CT and DBT, v is the vector form of the voxels of the target, and \mathbf{A} is the system matrix calculated by the Siddon algorithm.

$$P_\theta = \mathbf{A} \cdot v \quad (2.14)$$

To build up a system matrix (\mathbf{A}) using the Siddon algorithm, we created a $2 \times 2 \times 2$ -size voxel cube model (8 voxels) by assigning $N_x = 2$, $N_y = 2$, and $N_z = 2$ to analyze the calculated alpha-parametric set. Then we set a central ray starting point from $P_1(0,0,7)$ to $P_2(0,0,-2)$ to scan $2 \times 2 \times 2$ -size voxel cube. We also rotated the ray with -25° and $+25^\circ$ around the rotating point $(0,0,0)$ which is the center of the cube model, thus obtained three projections for data acquisitions of the cube. The visualization of 3-projected system is given in Figure 2.

We calculated and sorted in ascending order the alpha-parametric set for the $2 \times 2 \times 2$ model applying the Siddon algorithm. However due to the creation of the same values after performing 2.8, in the length calculation (2.9), the term $\alpha(m) - \alpha(m - 1)$ produced 0 that also cause the 0 value of $l(m)$. To get rid of zero-value intersection length problem which causes data loss in the system matrix of 3D image reconstruction modality

as defined in 2.14, we applied a function after performing 2.8, for ensuring two consecutive elements in the alpha-parametric set were not having of the same value. The unique function in Matlab 2018b (license number: 40731897) was applied to the alpha set for eliminating the same values in order not to cause data loss. The flowchart of the proposed method is demonstrated in Figure 3.

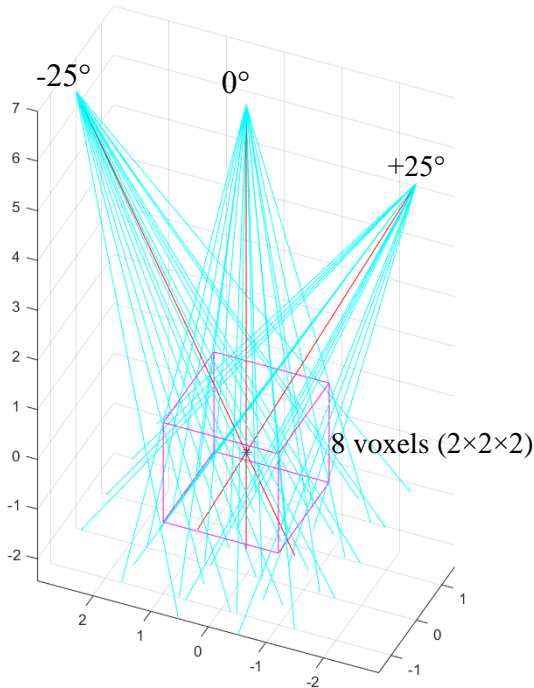


Figure 2. The visualization of the projection acquisition at $\theta = -25^\circ, 0^\circ, +25^\circ$ for $2 \times 2 \times 2$ -size (8 voxels) model of the Siddon algorithm

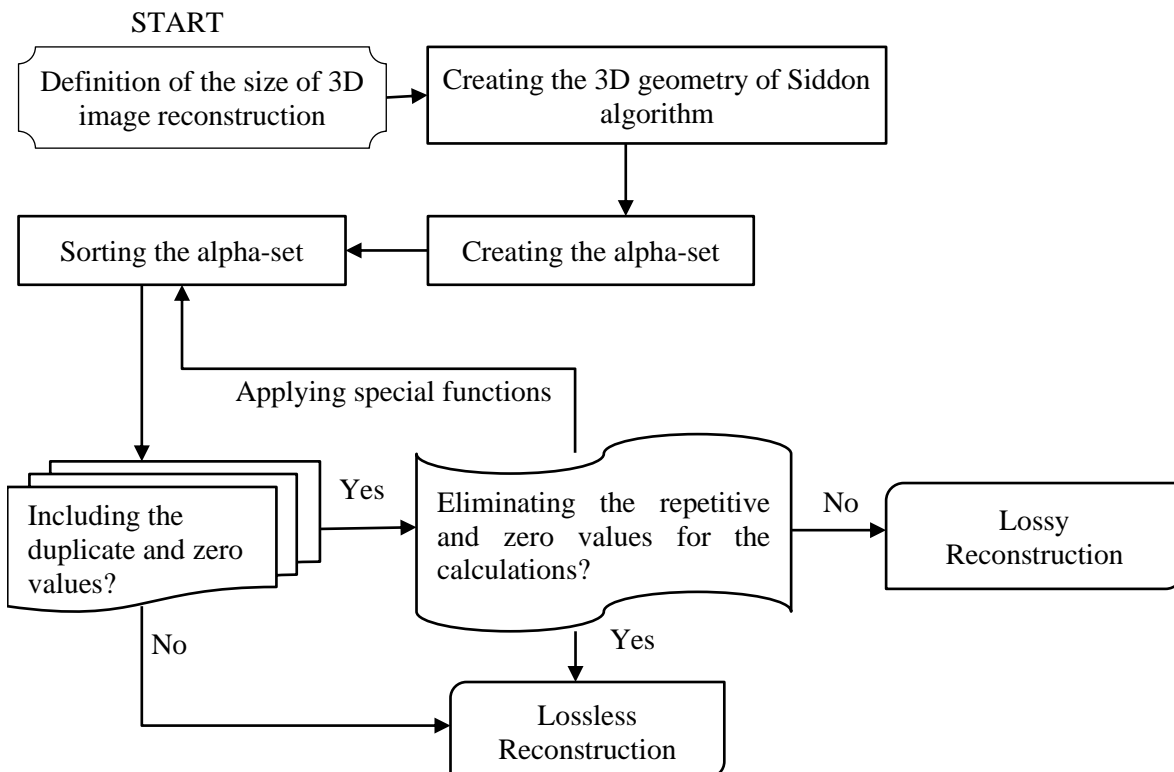


Figure 3. The flowchart of the proposed method

For a further and detailed analysis of data loss within the scope of the investigation of the alpha-parametric set, we 3D-reconstructed the images (with the size of $100 \times 100 \times 50$) of a synthetic multilayer phantom ($8 \text{ mm} \times 8 \text{ mm} \times 500 \mu\text{m}$) mimicking different microchannel (with $\sim 200 \mu\text{m}$ width and thickness) media separated with the various geometric structures (a circle, a square, and a diamond). For 3D image reconstruction, the 11 projection images (2D: 100×100) of the synthetic phantom obtained from the mini-Opto tomography platform (Polat et al., 2019b) in the range of $\pm 25^\circ$ were used. The representation of the reference projection (100×100) of the phantom both in the mini-Opto tomography platform imager and in Matlab 2018b is given in Figure 4.

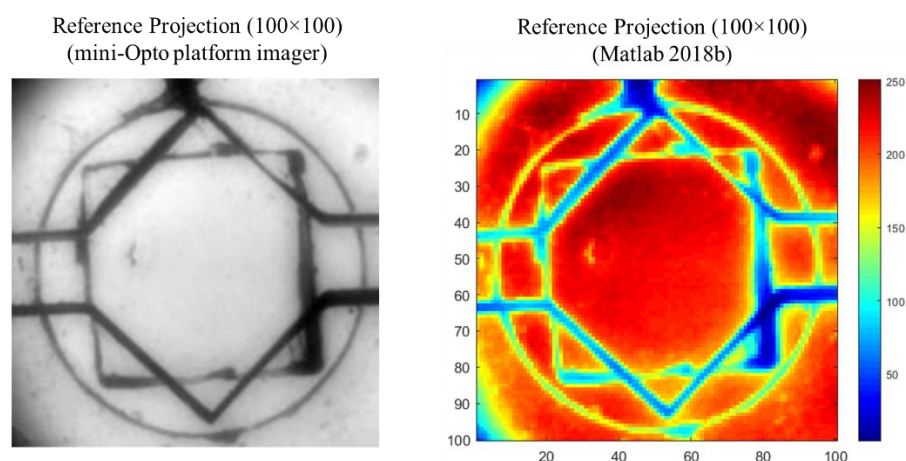


Figure 4. The representation of the reference projection (100×100) of the synthetic multilayer phantom mimicking different microchannel media separated with various geometric structures as a circle, a square, and a diamond both in the mini-Opto tomography platform imager (left) and in Matlab 2018b (right)

3. Results and Discussion

According to the acquisition geometry ($2 \times 2 \times 2$) described in the section of Materials and Methods, alpha-parametric set was calculated using (2.8) for the projections of -25° , 0° , and $+25^\circ$, and is given in the column of sorted alpha values without the unique function (column-2) in Table 1. It is clear that the repetitive values (0.6667, 0.8889, 2.0 for 6th ray, 0.6667, 0.8889 for 7th and 10th rays, 0.0, 0.6667, 0.8889 for 11th ray) that cause data loss in the system matrix were created in the set (system matrix without the unique function column in Table 2). The values of voxels of 1, 2, 3, and 4 at rays 6, 7, 10, and 11, respectively, shown in the column of system matrix without the unique function (8 voxels) were 0.0, which means data lost. Moreover, this situation may lead to huge data loss in 3D image reconstructions in medical imaging especially for huge dimensions such as size up to $800 \times 800 \times 50$. However, using the unique function, we eliminated the same values from the set so that the system matrix conserves the data, which is used in 3D image reconstruction. Additionally, using (2.5) the range alpha-set is limited between the values of min-alpha (0.6667) and max-alpha (0.8889), thus the values of 0.0 and 2.0 were also eliminated from final alpha set which is given in the last column, sorted alpha values with the unique function, of Table 1. Thanks to this result, the values (1.003) of voxels of 1, 2, 3, and 4 at rays 6, 7, 10, and 11, respectively, shown in the column of system matrix with the unique function (8 voxels) in Table 2, were created and data loss was prevented.

Table 1

Sorted alpha-set with and without the unique function at the projections of -25° , 0° , and $+25^\circ$

Ray values	Sorted alpha values without the unique function	Min-alpha	Max-alpha	Sorted alpha values with the unique function
6	0.6667, 0.6667, 0.7778, 0.8889, 0.8889, 2.0, 2.0	0.6667	0.8889	0.6667, 0.7778, 0.8889
7	0.0, 0.6667, 0.6667, 0.7778, 0.8889, 0.8889, 2.0	0.6667	0.8889	0.6667, 0.7778, 0.8889
10	0.0, 0.6667, 0.6667, 0.7778, 0.8889, 0.8889, 2.0	0.6667	0.8889	0.6667, 0.7778, 0.8889
11	0.0, 0.0, 0.6667, 0.6667, 0.7778, 0.8889, 0.8889	0.6667	0.8889	0.6667, 0.7778, 0.8889

Table 2
System matrix with and without the unique function at the projections of -25° , 0° , and $+25^\circ$

Ray	System matrix without the unique function (8 voxels)							
1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
6	0.0	0.0	0.0	0.0	1.003	0.0	0.0	0.0
7	0.0	0.0	0.0	0.0	0.0	0.0	1.003	0.0
8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
9	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
10	0.0	0.0	0.0	0.0	0.0	1.003	0.0	0.0
11	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.003
Ray	System matrix with the unique function (8 voxels)							
1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
6	1.003	0.0	0.0	0.0	1.003	0.0	0.0	0.0
7	0.0	0.0	1.003	0.0	0.0	0.0	1.003	0.0
8	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
9	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
10	0.0	1.003	0.0	0.0	0.0	1.003	0.0	0.0
11	0.0	0.0	0.0	1.003	0.0	0.0	0.0	1.003

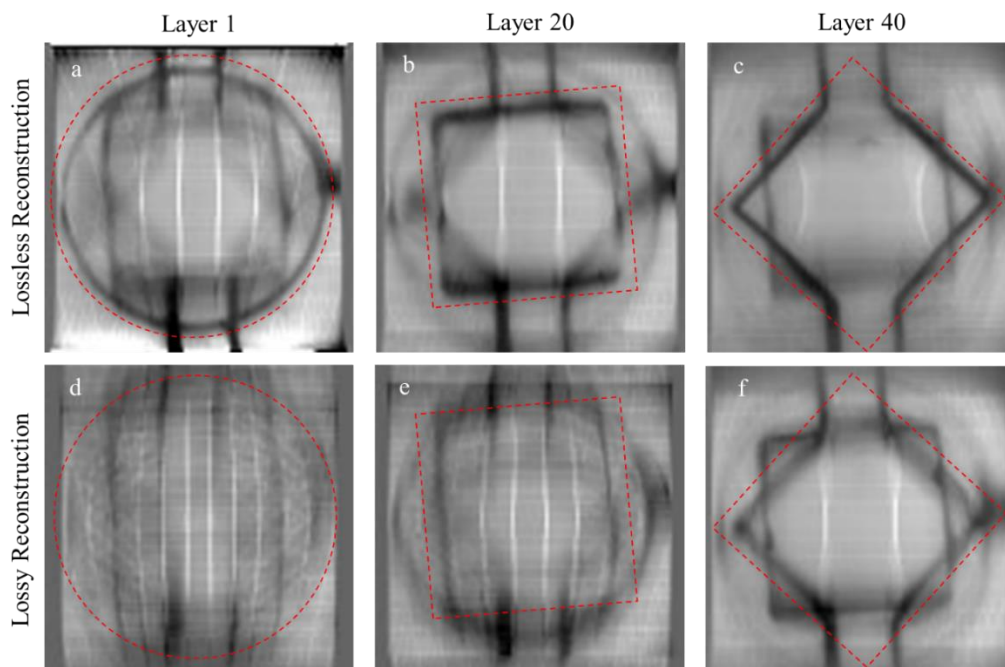


Figure 5. Lossless (first row: a, b, and c) and lossy (second row: d, e, and f) reconstruction. The geometric structures of the circle (a), the square (b), and the diamond (c) were captured in the layers of 1 (a), 20 (b), and 40 (c), respectively applying system matrix with the unique function (lossless reconstruction). The geometric structures of the circle (d), the square (e), and the diamond (f) could not captured in the layers of 1 (d), 20 (e), and 40 (f), respectively applying system matrix without the unique function (lossy reconstruction)

In this study, we tested and validated the effects of alpha-set on reconstructed images of the synthetic multilayer phantom that includes the geometric structures of the circle, the square, and the diamond with the size of $100 \times 100 \times 50$ (Figure 5). When Figure 5 is examined, the first row (a, b, and c) are the lossless reconstruction, which means the system matrix was implemented with the unique function. However, the second row (d, e, and f) are the lossy reconstruction, which means the system matrix was implemented without the unique function. In the qualitative assessment, it is clear that the circle, the square, and the diamond shapes in the lossless reconstruction were captured in the layers of 1, 20, and 40, respectively. Applying special functions in Matlab 2018b (e.i. unique function), the repetitive and zero-values of the alpha-parametric set were eliminated for the ray path calculations and lossless reconstruction. Therefore, the importance and the effect of alpha-set analysis in Siddon algorithm on reconstructed images should be taken into consideration in medical imaging modalities to avoid lossy reconstruction.

Evaluating the calculation values of alpha set and system matrix, we strongly recommend using the unique function after (2.8) step in the Siddon algorithm to avoid loss in data in system matrix for the 3D image reconstruction modality.

4. Conclusion

The Siddon algorithm provides an efficient radiological path calculation for building up the system matrix which is used in 3D image reconstruction applications such as computed tomography (CT) and digital breast tomosynthesis (DBT) modalities. The system matrix includes the voxel array and intersection length array for the interaction of the ray with the target object and is used to reconstruct the projections acquired from the target for 3D imaging. Thus, obtaining correct data in the system matrix by the Siddon algorithm is a very critical issue. The value of the elements of the system matrix is calculated by performing the calculation step of alpha-set parameters in the Siddon algorithm. Due to appearing the consecutive values of some elements, the intersection length calculation produces zero values that cause data loss in 3D image reconstruction. In this study, we proposed an additional step in order to eliminate the same values of the consecutive values of the alpha-parameter set which cause zero-valued intersection lengths. We performed both the Siddon algorithm in the regular process and the Siddon algorithm with the additional step we proposed to avoid zero-valued of the intersection length. Comparing the results in the system matrix for both methods, our proposal for the Siddon algorithm provided not produce zero-valued of intersection length, thus avoiding the data loss in the system matrix which is used in 3D image reconstructions in medical imaging. We proved our proposal by validating the concept with the lossy and lossless 3D image reconstruction of a $100 \times 100 \times 50$ size-synthetic phantom. For future work, we aim to perform and test our proposal for the Siddon algorithm for the increased dimensions of the system and applying it to the realistic phantoms

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Author Contributions

Adem Polat: Conceived the original concept, designed the analysis and wrote the manuscript.

Conflicts of Interest

The authors declare no conflict of interest.

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