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► To cite this version:

Addoum Ahmad, Olivier Farges, Fatmir Asllanaj. Optical properties reconstruction using the adjoint method based on the radiative transfer equation. *Journal of Quantitative Spectroscopy and Radiative Transfer*, Elsevier, 2018, pp.179-189. <10.1016/j.jqsrt.2017.09.015>. <hal-01610131>

HAL Id: hal-01610131

<https://hal.archives-ouvertes.fr/hal-01610131>

Submitted on 26 Jan 2018

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Optical properties reconstruction using the adjoint method based on the radiative transfer equation

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Abstract

An efficient algorithm is proposed to reconstruct the spatial distribution of optical properties in heterogeneous media like biological tissues. The light transport through such media is accurately described by the radiative transfer equation in the frequency-domain. The adjoint method is used to efficiently compute the objective function gradient with respect to optical parameters. Numerical tests show that the algorithm is accurate and robust to retrieve simultaneously the absorption μ_a and scattering μ_s coefficients for lowly and highly absorbing medium. Moreover, the simultaneous reconstruction of μ_s and the anisotropy factor g of the Henyey-Greenstein phase function is achieved with a reasonable accuracy. The main novelty in this work is the reconstruction of g which might open the possibility to image this parameter in tissues as an additional contrast agent in optical tomography.

Keywords Optical properties; image reconstruction; radiative transfer equation; adjoint method; crosstalk problem; anisotropy factor.

1 Introduction

Diffuse Optical Tomography (DOT) is a non-invasive imaging modality which employs a visible or Near infrared Laser source for probing biological tissues and measures light intensities at the boundary surface. In recent years, potential applications of DOT have been developed such as breast cancer detection [1] and brain functional imaging [2]. This tech-

nique seeks to recover the spatial distributions of optical properties inside the medium through an image reconstruction algorithm. Optical properties are different between healthy and cancerous tissues [3] [4] due to physiological or pathological changes [5]. In spite of all the success of DOT in cancer diagnosis applications, to date, only absorption (μ_a) and reduced scattering (μ'_s) coefficients reconstruction can be found in the literature. However, the anisotropy factor g of the Henyey-Greenstein (H-G) phase function has an important effect on light propagation [6] and can reveal rich informations on the anisotropic scattering behavior of the tissue: [7] showed that the g value of porcine brain tissue increases from 0.561 to 0.834 after thermal coagulation, [8] demonstrated that the anisotropy factor g of rat liver decreases from 0.952 to 0.946 in a tumor at 633 nm, [9] proved experimentally that g was different for normal human liver tissue and liver metastases at three different wavelengths. That means that g can also be modified when tissue is affected by an eventual tumor besides μ_a and μ_s . Therefore, this factor can provide a potential contrast agent for optical medical diagnosis between healthy and tumoral tissues. To our best knowledge, up to now, no works have been done to investigate the reconstruction of the spatial distribution of g [10]. On the other hand, an efficient forward model to predict light transport in the biological tissue is required in order to estimate optical parameters. Many research groups have adopted the diffusion equation as a forward model [11, 12, 13]. However, this model fails to accurately predict light propagation close to sources and boundaries, and also in highly absorbing mediums [14]. An additional drawback is that the estimation of g is not possible (g is assumed to be constant and known), due to the introduction of the reduced scattering coefficient and then the loss of information about the scattering phase function [6]. To overcome these limitations, more and more interest is turned towards a forward model based on the radiative transfer equation (RTE). The RTE rigorously describes the light propagation in biological tissues. The anisotropy factor g is an independant parameter in the RTE via the H-G phase function. Different forms of the RTE have been used in DOT. The frequency-domain (FD) approach [15, 16, 17, 18, 19, 20, 21] is the most widely employed, since it is a good trade-off between time-domain [22, 23, 24] and steady-state domain [25]. Moreover, the FD approach provides additional information (phase shift) compared to the steady-state modality and avoids the technical limitations of the experimental setup for time-domain often expensive. In addition, the use of FD data allows to better separate the optical properties than the steady-state data by reducing the crosstalk issue when simultaneous estimation is applied [16]. Another challenging task still remains in DOT: the inverse problem. The inversion algorithm can be considered as a large-scale optimization problem, since the optical properties vary spatially inside the medium. In principle, the simultaneous estimation of the three optical properties (μ_a , μ_s and g) is not possible, due to the non-uniqueness of the ill-posed problem when several optical properties distributions lead to an identical set of boundary data [6]. Additionally, these three parameters differ in nature, units, order of magnitude and sensitivities on the emerging intensity of the forward

model which makes the estimation inextricable. That's why, in this work, we reconstructed only two parameters either (μ_a and μ_s) or (μ_s and g) simultaneously in order to reduce the ill-posed nature of the problem. This inversion aims at recovering the optical properties of tissue through the minimization of an appropriate objective function (OF). Most of the time, the OF is the least-square error norm between the measured and the predicted data calculated by the forward model. Gradient-based algorithms are commonly used as optimization methods [13, 16, 26], which employ the gradient of the OF with respect to the optical parameters to find the minimum. These methods proved to be efficient and robust in DOT [24]. On the other side, the core problem and difficulty in the inversion procedure is to accurately compute the gradient of the OF. This process can be computational-intensive due to the number of parameters to retrieve which are space dependent. Generally, the adjoint differentiation is the most commonly used method for calculating the gradient because it uses only elementary results at each iteration step of the forward model [27, 28]. However, when the dimension of the problem is high (larger than 1000 for example), the use of this method becomes cumbersome and computationally expensive. More recently, [29] employed the adjoint method which gives an efficient and fast way to compute the OF gradient regardless of the number of unknowns. This is done by solving an additional (adjoint) equation for the adjoint variable whose computational cost is equivalent to that of the forward calculation.

In this work, a gradient-based algorithm using the RTE as forward model is employed to reconstruct the optical properties (μ_a , μ_s and g) of a heterogeneous medium. The gradient of the OF is obtained accurately by means of the adjoint method in the FD. The objective of this study is to test the efficiency and the robustness of the proposed algorithm in presence of some issues encountered in the DOT. These issues such as the collimated source number, the crosstalk between two optical parameters, the inclusion contrast level, the highly absorbing medium, the measurements noise level and the inclusion location effects are examined through several test cases. Furthermore, for author's best knowledge, the estimation of g and the simultaneous reconstruction of μ_s and g have not been reported yet in the previous works. This explains our motivation to test in particular the feasibility of the present method to reconstruct simultaneously μ_s and g with and without crosstalk. First, the RTE equations are described and the detector predictions on the boundary are given. Second, the adjoint method is presented through a lagrangian formalism for the computation of the OF gradient at multiple modulation frequencies. Finally, to illustrate the performance of the algorithm, single and simultaneous reconstructions of optical properties based on numerical test phantoms are presented in presence of certain issues mentioned above.

2 Forward model

In DOT, the light transport in the biological tissues is a forward model which aims at computing the prediction of the detectors reading once the source and the optical properties of the medium are known. The biological tissue is illuminated by an external collimated Laser beam $\Upsilon(\mathbf{r}_s, \omega_k)$ at the source position \mathbf{r}_s on the surface. ω_k is the modulation frequency of the intensity-modulated Laser source. In order to take into account this collimated light, the energy arriving in the medium is separated into two components $\psi = \psi_c + \psi_s$, respectively the collimated ψ_c and scattered radiance ψ_s . The ψ_c radiance is governed by the RTE state equation \mathcal{R}_c in the collimated direction $\mathbf{\Omega}_c$ and is solved analytically.

$$\mathcal{R}_c = \left[\mathbf{\Omega}_c \cdot \nabla + \left(\frac{i \omega_k}{v} + \mu_t(\mathbf{r}) \right) \right] \psi_c(\mathbf{r}, \omega_k) = 0. \quad (1)$$

The velocity of light, v , in tissue is the ratio $v = c/n$ of the velocity of light in vacuum and the refractive index of tissue. The total extinction coefficient $\mu_t(\mathbf{r})$, at position \mathbf{r} , is the sum of the absorption $\mu_a(\mathbf{r})$ and the scattering $\mu_s(\mathbf{r})$ coefficients. The boundary condition for the collimated component $\psi_c(\mathbf{r}, \omega_k)$ is given by:

$$\psi_c(\mathbf{r}, \omega_k) = \Upsilon(\mathbf{r}_s, \omega_k) \quad \text{for} \quad \mathbf{\Omega}_c \cdot \mathbf{n} < 0, \quad (2)$$

where \mathbf{n} is the outward normal unit vector of the boundary. It should be noted that the component Υ , in Eq. 2, represents only the transmitted part (no reflexion) of the collimated Laser beam into the medium. The scattered radiance $\psi_s(\mathbf{r}, \mathbf{\Omega}, \omega_k)$ is obtained by solving the RTE state equation \mathcal{R}_s in the direction $\mathbf{\Omega}$ of the light propagation such as :

$$\mathcal{R}_s = \left[\mathbf{\Omega} \cdot \nabla + \left(\frac{i \omega_k}{v} + \mu_t(\mathbf{r}) \right) \right] \psi_s(\mathbf{r}, \mathbf{\Omega}, \omega_k) - \mu_s(\mathbf{r}) \int_{\Omega'=2\pi} p(\mathbf{\Omega}', \mathbf{\Omega}) \psi_s(\mathbf{r}, \mathbf{\Omega}', \omega_k) d\Omega' - S_c(\mathbf{r}, \mathbf{\Omega}, \omega_k) = 0. \quad (3)$$

The H-G phase function $p(\mathbf{\Omega}' \cdot \mathbf{\Omega})$ is the most widely adopted scattering phase function in biomedical optics and has been used here [23, 30]. This function, is the probability that photons traveling in direction $\mathbf{\Omega}'$ scatter into direction $\mathbf{\Omega}$. The H-G phase function mathematical expression in 2D is given by:

$$p(\mathbf{\Omega}' \cdot \mathbf{\Omega}) = \frac{1}{2\pi} \frac{1 - g^2(\mathbf{r})}{(1 + g^2(\mathbf{r}) - 2g(\mathbf{r}) \mathbf{\Omega}' \cdot \mathbf{\Omega})}. \quad (4)$$

The anisotropy factor $g(\mathbf{r})$ represents the mean cosine of the angles of the scattered directions $\mathbf{\Omega}$ with respect to the incident ones $\mathbf{\Omega}'$. This factor is spatially dependent in our case for the heterogeneous medium. The source term $S_c(\mathbf{r}, \mathbf{\Omega}, \omega_k)$ in Eq. (3) induced by the scattering of the collimated radiance $\psi_c(\mathbf{r}, \omega_k)$ is given by:

$$S_c(\mathbf{r}, \mathbf{\Omega}, \omega_k) = \mu_s(\mathbf{r}) p(\mathbf{\Omega}_c, \mathbf{\Omega}) \psi_c(\mathbf{r}, \omega_k). \quad (5)$$

Eq. (3) is associated to a semi-transparent boundary condition [31] with Fresnel reflection at the interface (air / biological tissue) due to the refractive index mismatch. The detector prediction $P(\mathbf{r}_d, \omega_k)$ corresponding to the exitance at the

detector position \mathbf{r}_d on the illuminated surface is obtained by:

$$P(\mathbf{r}_d, \omega_k) = \int_{\mathbf{n} \cdot \boldsymbol{\Omega}' > 0} [1 - \rho(\Theta)] \psi_s(\mathbf{r}, \boldsymbol{\Omega}', \omega_k) (\boldsymbol{\Omega}' \cdot \mathbf{n}) d\Omega', \quad (6)$$

where $\rho(\Theta)$ is the reflectivity of the surface $\partial\mathcal{D}$. The forward model has been solved accurately with a Modified Finite Volume Method (MFVM). The methodology of this method is not repeated here and we refer the reader to [32], for details. The stability and accuracy of the MFVM have been validated through comparisons with the Monte Carlo (MC) technique and analytical solution of RTE on available test cases. The MFVM, compared to other deterministic numerical solutions of the RTE (available in the literature) has the advantage to have a high precision with an error less than 1% with respect to MC simulations and RTE analytical solution. This is mainly due to the fact that the RTE is also solved inside each control volume through an exponential schema.

3 Inverse problem

In the following, we first define the discrete sum of the objective function over all modulation frequencies. Then we introduce the minimization problem where the state equations are used as constraints. After introducing the Lagrangian formalism, we show how to deduce the adjoint equations and the objective function gradient.

3.1 Objective function and minimization problem

The OF $\mathcal{J}(\beta)$ to be minimized in the inversion procedure is the mean square discrepancy between the measurements, $M(\mathbf{r}_d, \omega_k)$, and the predictions of the forward model based on the RTE, $P(\mathbf{r}_d, \omega_k)$, at detector positions, \mathbf{r}_d on the surface of the medium over all intensity modulation frequencies ω_k :

$$\mathcal{J}(\beta) = \frac{1}{2} \sum_{\omega_k=1}^{N_\omega} J(\beta, \omega_k) \quad \text{with} \quad J(\beta, \omega_k) = \sum_{d=1}^{N_d} \|P(\mathbf{r}_d, \omega_k) - M(\mathbf{r}_d, \omega_k)\|^2, \quad (7)$$

The vector β contains the spatial distribution of the optical properties in the heterogeneous medium. Here N_d and N_ω are the total numbers of detectors and modulation frequencies, respectively. The goal of the optimization technique is to determine the vector $\hat{\beta}$ that minimizes the OF iteratively. This vector $\hat{\beta}$ will be a solution to the minimization problem and is displayed as a two-dimensional optical image.

3.2 Lagrangian and adjoint model equations

The adjoint equations are derived by considering that the OF at each modulation frequency $J(\beta, \omega_k)$ must be minimized under some constraints given by the RTE state equations at the corresponding frequency ω_k . Hence, we can rewrite the

original OF by following the Lagrangian formalism of the minimization problem as:

$$\mathcal{L}(\beta, \psi_s, \psi_c, \lambda_s, \lambda_c) = \sum_{\omega_k=1}^{N_\omega} \left[J(\beta, \omega_k) + \left(\lambda_s \left| \mathcal{R}_s(\beta, \psi_s, \psi_c) \right|_s \right)^{\omega_k} + \left(\lambda_c \left| \mathcal{R}_c(\beta, \psi_c) \right|_c \right)^{\omega_k} \right], \quad (8)$$

where $\lambda_c = \lambda_c(\mathbf{r}, \omega_k)$, $\lambda_s = \lambda_s(\mathbf{r}, \mathbf{\Omega}, \omega_k)$ are the complex adjoint variables to ψ_c and ψ_s , respectively. The inner products $(\cdot|\cdot)_c^{\omega_k}$ and $(\cdot|\cdot)_s^{\omega_k}$ associated to the solution space, respectively, of ψ_c and ψ_s are defined by:

$$\left(\lambda_s \left| \mathcal{R}_s \right|_s \right)^{\omega_k} = \mathbf{Re} \int_{\mathcal{D}} \int_{\Omega=2\pi} \overline{\lambda_s(\mathbf{r}, \mathbf{\Omega}, \omega_k)} \mathcal{R}_s(\beta, \psi_s, \psi_c) d\Omega dr \quad (9)$$

$$\left(\lambda_c \left| \mathcal{R}_c \right|_c \right)^{\omega_k} = \mathbf{Re} \int_{\mathcal{D}} \overline{\lambda_c(\mathbf{r}, \omega_k)} \mathcal{R}_c(\beta, \psi_c) dr. \quad (10)$$

When ψ_c, ψ_s verify the state equations Eqs. (1), (3), respectively, that leads to:

$$\mathcal{L}(\beta, \psi_s, \psi_c, \lambda_s, \lambda_c) = \mathcal{J}(\beta), \quad \mathcal{L}'(\beta, \psi_s, \psi_c, \lambda_s, \lambda_c) = \mathcal{J}'(\beta), \quad (11)$$

Using Eq. (11), we can extract the gradient of the OF from the L_2 inner product of the directional differential \mathcal{L}' with respect to β in the direction $\delta\beta$, such that:

$$\mathcal{L}'(\beta) = \left(\nabla \mathcal{J}(\beta) \left| \delta\beta \right|_{L_2} \right) \quad (12)$$

Notice that the functional \mathcal{L} is independant of λ_s and λ_c due to the fact that the residuals \mathcal{R}_s and \mathcal{R}_c are zero, yielding:

$$\frac{\partial \mathcal{L}(\beta, \psi_s, \psi_c)}{\partial \lambda_s} = 0, \quad \frac{\partial \mathcal{L}(\beta, \psi_s, \psi_c)}{\partial \lambda_c} = 0. \quad (13)$$

The lagrangian formalism assumes that the variation of $\partial \mathcal{L}$ is not non-zero unless there is a variation of β . This condition is ensured by a particular choice of adjoint variables λ_s and λ_c which allows to compute the OF gradient without having to compute the sensitivities $\delta\psi_s = (\partial\psi_s(\mathbf{r}, \mathbf{\Omega}, \omega_k)/\partial\beta)\delta\beta$ and $\delta\psi_c = (\partial\psi_c(\mathbf{r}, \omega_k)/\partial\beta)\delta\beta$. These sensitivities are computationally expensive. Hence, that leads to the following adjoint equations model of the FD DOT problem:

$$\partial_{\psi_s} \mathcal{L}(\beta, \psi_s, \psi_c) \delta\psi_s = 0, \quad \partial_{\psi_c} \mathcal{L}(\beta, \psi_s, \psi_c) \delta\psi_c = 0. \quad (14)$$

The adjoint equations can be obtained by partially differentiating the Lagrangian functional Eq. (8) with respect to ψ_s, ψ_c in direction $\delta\psi_s$ and $\delta\psi_c$, respectively. After using the definition of the adjoint operator [33] and the inner products properties, we can reformulate the adjoint equations over N_ω modulation frequencies ω_k such as:

$$\partial_{\psi_s} \mathcal{L}(\beta, \psi_s, \psi_c) \delta\psi_s = \sum_{\omega_k=1}^{N_\omega} \left[\frac{\partial J(\beta, \omega_k)}{\partial \psi_s} + \left(\frac{\partial \mathcal{R}_s(\beta, \psi_s, \psi_c)}{\partial \psi_s} \right)^* \lambda_s \right] = 0, \quad (15)$$

$$\partial_{\psi_c} \mathcal{L}(\beta, \psi_s, \psi_c) \delta\psi_c = \sum_{\omega_k=1}^{N_\omega} \left[\left(\frac{\partial \mathcal{R}_s(\beta, \psi_s, \psi_c)}{\partial \psi_c} \right)^* \lambda_s + \left(\frac{\partial \mathcal{R}_c(\beta, \psi_c)}{\partial \psi_c} \right)^* \lambda_c \right] = 0. \quad (16)$$

Note that $\partial J(\beta, \omega_k)/\partial \psi_c = 0$, since the OF is independent on the collimated light source.

By using the definitions in Eq. (1) and (3) for \mathcal{R}_c and \mathcal{R}_s , respectively, the adjoint variables must be a solution of the following adjoint equations system at each modulation frequency ω_k :

$$\left[-\mathbf{\Omega} \cdot \nabla + \left(\frac{-i \omega_k}{v} + \mu_t(\mathbf{r}) \right) \right] \lambda_s(\mathbf{r}, \mathbf{\Omega}, \omega_k) - \mu_s(\mathbf{r}) \int_{\Omega'=2\pi} p(\mathbf{\Omega}', \mathbf{\Omega}) \lambda_s(\mathbf{r}, \mathbf{\Omega}', \omega_k) d\mathbf{\Omega}' + \frac{\partial J(\beta, \omega_k)}{\partial \psi_s} = 0 \quad (17)$$

$$\left[-\mathbf{\Omega}_c \cdot \nabla + \left(\frac{-i \omega_k}{v} + \mu_t(\mathbf{r}) \right) \right] \lambda_c(\mathbf{r}, \omega_k) - \mu_s(\mathbf{r}) \int_{\Omega=2\pi} p(\mathbf{\Omega}_c, \mathbf{\Omega}) \lambda_s(\mathbf{r}, \mathbf{\Omega}, \omega_k) d\mathbf{\Omega} = 0. \quad (18)$$

The system of the RTE adjoint equations can be solved with the same solution method as the forward model equations.

In the next section, we will show how to obtain the gradient vector components $\nabla \mathcal{J}$ with respect to the optical properties (μ_a , μ_s and g) by simply using the adjoint variables $\lambda_s(\mathbf{r}, \mathbf{\Omega}, \omega_k)$ and $\lambda_c(\mathbf{r}, \omega_k)$ of the adjoint model described above.

3.3 Gradient expressions

The directional derivative of the objective function \mathcal{J}' is equal to that of the lagrangian functional \mathcal{L}' (see Eq. (11)). Thus, it suffices to determine the latter in order to extract the gradient $\nabla \mathcal{J}$. By applying Eqs. (11 - 14), the gradient $\nabla \mathcal{J}$ can be deduced by differentiating the lagrangian functional \mathcal{L} with respect to β in direction $\delta\beta$ such as:

$$\left(\nabla \mathcal{J}(\beta) \middle| \delta\beta \right)_{L_2} = \sum_{\omega_k=1}^{N_\omega} \left[\left(\lambda_c \middle| \frac{\partial \mathcal{R}_c(\beta, \psi_c)}{\partial \beta} \delta\beta \right)_c^{\omega_k} + \left(\lambda_s \middle| \frac{\partial \mathcal{R}_s(\beta, \psi_s, \psi_c)}{\partial \beta} \delta\beta \right)_s^{\omega_k} \right]. \quad (19)$$

Note that $\partial J(\beta, \omega_k)/\partial \beta = 0$ as the OF does not depend explicitly on β (see Eq.(7)).

This latter expression clearly shows that only a simple inner product has to be calculated. It should be noted that, the gradient of the OF at multiple frequencies is the sum of all the gradients computed at each modulation frequency ω_k .

$$\left(\nabla \mathcal{J}(\beta) \middle| \delta\beta \right)_{L_2} = \sum_{\omega_k=1}^{N_\omega} \left(\nabla J(\beta, \omega_k) \middle| \delta\beta \right)_{L_2}. \quad (20)$$

Applying Eq. (19) for $\delta\mu_a$, $\delta\mu_s$ and δg , we obtain an analytical expression of the OF gradient, with respect to μ_a , μ_s and g , respectively:

$$\left(\nabla \mathcal{J}(\mu_a) \middle| \delta\mu_a \right)_{L_2} = \sum_{\omega_k=1}^{N_\omega} \left[\left(\lambda_s \middle| \psi_s \delta\mu_a \right)_s^{\omega_k} + \left(\lambda_c \middle| \psi_c \delta\mu_a \right)_c^{\omega_k} \right]. \quad (21)$$

$$\begin{aligned} \left(\nabla \mathcal{J}(\mu_s) \middle| \delta\mu_s \right)_{L_2} &= \sum_{\omega_k=1}^{N_\omega} \left[\left(\lambda_s \middle| \psi_s \delta\mu_s \right)_s^{\omega_k} + \left(\lambda_c \middle| \psi_c \delta\mu_s \right)_c^{\omega_k} \right. \\ &\quad \left. - \left(\lambda_s \middle| \left(\int_{\Omega'=2\pi} \psi_s p(\mathbf{\Omega}', \mathbf{\Omega}) d\mathbf{\Omega}' + \psi_c p(\mathbf{\Omega}_c, \mathbf{\Omega}_k) \right) \delta\mu_s \right)_s^{\omega_k} \right] \end{aligned} \quad (22)$$

$$\left(\nabla \mathcal{J}(g) \middle| \delta g \right)_{L_2} = \sum_{\omega_k=1}^{N_\omega} \left(\lambda_s \middle| -\mu_s \left(\int_{\Omega'=2\pi} \psi_s \frac{\partial p(\mathbf{\Omega}', \mathbf{\Omega})}{\partial g} d\mathbf{\Omega}' + \psi_c \frac{\partial p(\mathbf{\Omega}_c, \mathbf{\Omega})}{\partial g} \right) \delta g \right)_s^{\omega_k} \quad (23)$$

The gradient is then computed for all the optical parameters regardless of the number of unknowns of the problem.

Thus, the adjoint formulation gives a fast way to efficiently compute the gradient. This is done by solving an additional

equation for the adjoint variables and then evaluating the gradient through a simple inner product. To our knowledge, the gradient expression $\nabla \mathcal{J}(g)$ with respect to g is derived for the first time by using the adjoint model of the RTE. Once the gradient $\nabla \mathcal{J}(\beta)$ is obtained accurately, a reconstruction scheme based on the *Limited memory* (Lm) BFGS method [29] is employed in order to update the spatial distribution of the optical properties. The advantage of this method is that it avoids the cost of building the Hessian and inverting it as in the Newton methods [34, 35]. After computing the Lm-BFGS search direction, the *Armijo* line search [36] is employed to find an optimal step size, α^k , which minimises sufficiently the OF.

4 Results and discussions

4.1 Model description

The reconstruction of the spatial distribution of optical properties in a non-homogeneous biological medium is studied. A two-dimensional $4 \times 4 \text{ mm}^2$ domain which contains three circular inclusions is examined as shown in Fig. 1(a). The inclusions (0.5 mm in diameter each, centered at $x = -0.1 \text{ mm}$; $y = 1 \text{ mm}$ for inclusion A, $x = 1 \text{ mm}$; $y = 1 \text{ mm}$ for inclusion B and $x = -1 \text{ mm}$; $y = -1 \text{ mm}$ for inclusion C) are embedded as heterogeneities in the background medium.

[Figure 1 about here.]

The optical properties of these inclusions can take different values for the different test cases considered below. The refractive index of the medium is uniformly set at $n = 1.4$ while that of the surrounding medium (air) is set to unity. For all these cases, the homogeneous background optical properties are used as the initial guesses to start the inverse procedure. Four Gaussian Laser sources illuminate simultaneously the mid-center of each side of the medium, unless the cases specified otherwise (those using one source). The expression of the Gaussian function in space along x -axis or y -axis ($s = x$ or y) is given by:

$$\Upsilon(s) = \frac{1}{\sigma_s \sqrt{2\pi}} \exp\left(-\frac{s^2}{2\sigma_s^2}\right), \quad (24)$$

where $\sigma_s = 0.5 \text{ mm}$ is the standard deviation of the spatial Gaussian beam. Eighty detectors are distributed around the numerical phantom (one source and 20 detectors are located on each side of the medium). All detectors predictions were used in the minimization problem, except the cases where μ_s and g factor are reconstructed simultaneously. The synthetic data were obtained by running the forward model using the exact heterogeneous distribution of the optical properties we want to reconstruct. An unstructured triangular mesh of $N_s = 2577$ nodes (degrees of freedom) corresponding to 4992 triangles was used (Fig. 1(b)). The angular space was discretized into 32 directions and each direction was also subdivided

into 8 solid angles for the normalized phase function. The reconstructions were achieved by fitting the FD data obtained at 10 modulation frequencies equally distributed in the range of 100 MHz to 1 GHz. We have found that the multifrequency approach provides a better estimation quality than that at single modulation frequency [37, 38, 39, 40]. This is because the underdetermination of the large-scale optimization problem was reduced by employing multiple modulation frequencies. The reconstruction process is terminated when the normalized difference of the OF between two subsequent iteration steps was smaller than $\epsilon = 10^{-4}$. To compare the quality of reconstruction images, the relative error ε between real and reconstructed values of optical properties is defined as:

$$\varepsilon_{\beta}(ZI) = \frac{100}{P} \sum_{i=1}^{ZI} \left\| \frac{\hat{\beta}_i - \beta_i^*}{\beta_i^*} \right\|, \quad (25)$$

where $\hat{\beta}_i$ and β_i^* are the reconstructed and the exact values of the optical parameter at the i^{th} node of the mesh, respectively. P represents the unknowns number (mesh nodes) in the zone of interest ZI of the image. This ZI can be either the background, inclusion or cross-talk zone or even the whole reconstructed image.

4.2 Impact of source number on reconstruction

As the first test problem, the effect of source number on reconstruction is examined. For this purpose, we consider the phantom described above where only the scattering coefficient varies spatially inside the medium through the inclusions. The optical properties of the background are chosen in the range of biological tissues ($\mu_a = 0.05 \text{ mm}^{-1}$, $\mu_s = 5 \text{ mm}^{-1}$ and $g = 0.9$). These properties yield to a highly-forward anisotropically scattering medium. Inclusions A and C are assigned the real scattering coefficient $\mu_s^* = 4 \text{ mm}^{-1}$, while inclusion B is assigned the real $\mu_s^* = 6 \text{ mm}^{-1}$, corresponding to a 20% decrease and increase, respectively, relative to the scattering coefficient of the background medium. Figs. 2(a) and 2(b) display the reconstructed images of μ_s when only the top surface of the medium was probed and when the four different boundaries of the medium were illuminated, respectively.

[Figure 2 about here.]

From Figs. 2(a,b), it can be seen that the three inclusions are spatially well recovered in the exact locations for both cases. As expected, the two top inclusions (A and B) are accurately reconstructed while the deeper one (inclusion C) is significantly overestimated ($\hat{\mu}_s = 4.5 \text{ mm}^{-1}$) when probing only the north boundary. In addition, the inclusion C has worse contrast and circular shape (Fig. 2(a)). However, the circular shape and the contrast of the inclusion C are clearly improved and enhanced when scanning the medium from all its boundaries. The scattering values are accurately estimated $\hat{\mu}_s = 4 \text{ mm}^{-1}$, $\hat{\mu}_s = 6 \text{ mm}^{-1}$ and $\hat{\mu}_s = 4 \text{ mm}^{-1}$ at the centers of inclusions A, B and C, respectively (Fig. 2(b)).

4.3 Simultaneous reconstruction of μ_a and μ_s

In this section, the simultaneous reconstruction of the spatially dependent absorption and scattering distribution is examined with three different test cases. The anisotropy factor of the H-G phase function is kept constant $g = 0.9$. We reconsider the same phantom as described in Fig. 1, illuminated by four collimated Laser beams.

The most encountered issues and challenges in DOT are reported in this section in order to assess the robustness of the reconstruction algorithm. Firstly, the "crosstalk" problem is frequently encountered in practice when simultaneous estimation of several parameters is applied. This problem is due to the non-uniqueness of the DOT where many combinations of optical properties can lead to similar boundary data. For this purpose, we assumed that the inclusion A represents a heterogeneity with low in both absorption and scattering coefficients compared to the background optical properties. Whereas, inclusion B is highly scattering only while inclusion C is highly absorbing only. This configuration is intended to mimic a crosstalk in the three test media. Secondly, the "contrast" level represents the difference between the background (initial optical value) and the inclusion (exact optical value). Different contrasts may lead to different reconstruction results [41, 42]. In the 3 test cases, the three inclusions represent a contrast of 20% with respect to the background optical values except for case 2 where inclusions B and C represent a contrast of 40%. This case is chosen in order to evaluate the effect of this contrast on the crosstalk issue and the estimation quality. Two homogeneous backgrounds of different optical properties are employed. The first two cases are assigned the low-absorbing background medium of $(\mu_a = 0.05 \text{ mm}^{-1}, \mu_s = 5 \text{ mm}^{-1})$ while the third one consists of a very high-absorbing medium $(\mu_a = 1 \text{ mm}^{-1}, \mu_s = 5 \text{ mm}^{-1})$. This last case presents a situation in which the diffusion approximation is not valid. The exact optical properties of the inclusions for each case are listed in Table 1. Figure 3 displays the reconstructed μ_a and μ_s images for the 3 test mediums. The relative reconstruction errors of background, inclusions and crosstalk are given in Table 2.

[Table 1 about here.]

[Figure 3 about here.]

As shown in Fig. 3, the inclusions are accurately located in both optical parameters for all test cases. For the two low-absorbing media, perturbations such as edges artifacts are more remarkable in the absorption images. Also, the local values in the scattering maps are accurately retrieved while that of the absorption maps are underestimated. The purely absorbing inclusion C has no crosstalk impact on the μ_s images even with high contrast of 40% (see Figs. 3(b,d)). This is because the use of the FD data is expected to better separate between the two parameters. However, the crosstalk

phenomenon is only pronounced in absorption images. As shown in Figs. 3(a,c), the scattering inclusion B appeared as a false positive heterogeneity in the μ_a images for both cases. This behavior is mainly due to the different sensitivities of the RTE model which is much more sensitive to variations in μ_s than μ_a [43]. These parameters have somehow the same effect on the boundary data. In other words, a decrease in intensity could be caused by either an increase in absorption or in scattering [6], since both contribute to light extinction in tissue. Therefore, the low emerging intensities caused by the highly scattering inclusion B, have been analyzed as a highly inclusion in both scattering and absorption (crosstalk).

By comparing the cases 1 and 2, we find that the estimation errors of the inclusions B and C with 20% of contrast, $\varepsilon_{\mu_s}^{IncB} = 5.46\%$ and $\varepsilon_{\mu_a}^{IncC} = 9.16\%$, have been approximately doubled to $\varepsilon_{\mu_s}^{IncB} = 9.70\%$ and $\varepsilon_{\mu_a}^{IncC} = 17.7\%$, respectively, when the contrast level was increased about 40% (see Table 2). However, the inclusion A error was slightly increased to $\varepsilon_{\mu_a}^{IncA} = 10.95\%$ and $\varepsilon_{\mu_s}^{IncA} = 7.35\%$ because the contrast level for this inclusion has not changed for the two cases. In the case 2, perturbations are more pronounced in the absorption background ($\varepsilon_{\mu_a}^{Background} = 6.11\%$) and some artifacts are also remarkable in the scattering image ($\varepsilon_{\mu_s}^{Background} = 1.36\%$). Thus, it is interesting to note that the high inclusion contrast leads to more artifacts around the boundary while the low inclusion contrast shows better estimation quality. As shown in Figs. 3(a,c), the reconstructed absorbing inclusion C is readily visible when the contrast is up to 40%. This is because the influence of the last becomes more important on the emerging data with higher contrast. Furthermore, it is also seen that the contrast has some considerable effect on the crosstalk. The crosstalk error in the absorption map $\varepsilon_{\mu_a}^{Crosstalk} = 12.29\%$ has been doubled to $\varepsilon_{\mu_a}^{Crosstalk} = 24.49\%$ when the contrast of the inclusion B was increased from 20% to 40%, respectively. For the scattering map the crosstalk error $\varepsilon_{\mu_s}^{Crosstalk} = 0.17\%$ was slightly increased to $\varepsilon_{\mu_s}^{Crosstalk} = 0.27\%$, but remains low. Hence, a large contrast level leads to a worse estimation quality and further to a strong crosstalk effect in the absorption images. This has an important implication because the realistic contrast levels between tumor and normal tissue are believed to be in the range of the lower contrast levels [42].

For the test case 3, the reconstruction of optical properties is achieved with a reasonable accuracy thanks to the RTE based forward model. This result would not have been possible with the diffusion approximation, since it fails to predict accurately the light propagation in such medium. Comparing now the tests 1 and 3, we can observe that the reconstruction of the μ_a background becomes clearer with less perturbations leading to lower relative error of $\varepsilon_{\mu_a}^{Background} = 0.94\%$ in the highly-absorbing medium (Figs. 3(a,e)). In addition, the crosstalk error induced by the scattering inclusion B has been reduced about 50% ($\varepsilon_{\mu_a}^{Crosstalk} = 5.769\%$) in the μ_a image. The estimation errors of the inclusions A and C ($\varepsilon_{\mu_a}^{IncA} = 11.62\%$, $\varepsilon_{\mu_a}^{IncC} = 9.94\%$) are somewhat similars compared to errors obtained in case 1. This is because the contrast level of 20% was kept unchanged in the two test cases. For the μ_s image, the background is accurately recovered with small

error of $\varepsilon_{\mu_s}^{Background} = 0.60\%$. In contrary to case 1 and 2, the purely absorbing inclusion C has now a significant crosstalk effect on the μ_s image (Figs. 3(b,f)). Nevertheless, the crosstalk error is still relatively small ($\varepsilon_{\mu_s}^{Crosstalk} = 1.85\%$) in the μ_s image. The estimation errors of the inclusions A and B were increased to almost 50% $\varepsilon_{\mu_s}^{IncA} = 12.1\%$ and $\varepsilon_{\mu_s}^{IncB} = 12.6\%$ compared to the case 1, respectively. One can deduce here that the reconstruction quality of the μ_a image is improved (low crosstalk and background errors) while that of the μ_s image is worse (high crosstalk and inclusions errors). This improvement of μ_a can be explained by the highly absorption coefficient yielding to a comparable order of magnitude with the scattering coefficient. In this situation, the RTE model has a significant sensitivity to any small variation of μ_a (20%) when the absorption coefficient is very high. Therefore, the algorithm provides a better reconstruction quality for μ_a . On the other hand, the highly absorbing coefficient produces a high light extinction in the tissue which attenuates the multiple scattering of light and therefore to μ_s inaccuracy.

On the other hand, if the μ_s and μ_a original images contain each two inclusions where their exact values are both either high or low with respect to their initial values, the proposed algorithm was tested and has been proven to reconstruct accurately this example case.

[Table 2 about here.]

4.4 Reconstruction of g

In this case, the anisotropy factor g varies spatially inside the medium through the inclusions as represented in Fig. 1(a). The algorithm was used to reconstruct the spatial distribution of g when μ_a and μ_s are assumed to be known. The optical properties of the homogeneous background are $\mu_a = 0.05 \text{ mm}^{-1}$, $\mu_s = 5 \text{ mm}^{-1}$ and $g = 0.9$. The inclusions A and C are assigned the exact value $g^* = 0.85$, while the inclusion B is assumed to have higher forward-scattering value $g^* = 0.95$. In order to test the robustness of the algorithm, the synthetic data are corrupted by adding normal distributed random errors to the exact predictions such as:

$$M_d = P_d^* + \text{rand } \sigma_d, \quad (26)$$

where M_d and P_d^* are the measured value and the exact prediction at the d -th detector on the bounding surface, respectively. The rand function, in Eq. (26), generates random numbers with a normal Gaussian distribution. The noise level σ_d is defined as the standard deviation of the measured value at the d -th detector. Four examples of different noise levels present in the synthetic data (0%, 3%, 6% and 10%) were used in the reconstructions. The results are depicted in Figure 4 and the computational features of the reconstruction algorithm are given in Table 3.

[Figure 4 about here.]

For the noise-free example, the reconstructed g image is in good agreement with the original medium (Fig. 4(a)). Locations and circular shapes of the inclusions were clearly reconstructed. The retrieved local values at the inclusions centers were accurately estimated leading to low relative error of the inverted image $\varepsilon_g = 0.29\%$. Also, the background anisotropy factor is well recovered for all the cases. These results show that the gradient expression with respect to the anisotropy factor (Eq. 23) has been validated and accurately computed by the present adjoint method. For the noisy data examples, it is also seen that the algorithm can well detect and locate the inclusions inside the medium (Figs. 4(b, c, d)). However, the edges artifacts are more pronounced and the circular shape of inclusions is distorted as the noise level is found to increase. Additionally, the estimation error increases to $\varepsilon_g = 0.41\%$, $\varepsilon_g = 0.53\%$ and $\varepsilon_g = 0.72\%$ when the noise level increases to 3%, 6% and 10%, respectively (see Tab. 3). As expected, higher noise levels on the boundary data lead to quality image degradation. From Table 3, the optimization procedure reached the stopping criterion faster for a higher noise level. This is because the OF converges around a certain noisy value. This computational feature is often encountered for gradient based algorithms and has been reported in [10, 36]. It should be noted here that the estimation of g would have not been possible with the diffusion approximation. Since, the information about the anisotropy factor is lost by considering the reduced scattering coefficient $\mu'_s = \mu_s(1 - g)$ in the diffusion equation.

[Table 3 about here.]

4.5 Simultaneous reconstruction of μ_s and g

4.5.1 With crosstalk problem

In this section, the spatial distributions of the scattering coefficient μ_s and the anisotropy factor g are reconstructed simultaneously. The original phantom to reconstruct contains only the two top inclusions A and B located as depicted in Fig. 1(a). The medium was probed by one Gaussian source at the mid-center of the top surface and only the backscattered light (reflectance) on the illuminated boundary was used for reconstruction. The use of one source allows to highlight the inclusion location effect with respect to the source on the estimation and crosstalk qualities. In this source-detectors configuration, we are able to assess the sensitivities of the μ_s coefficient and the g factor on the reflectance. The homogeneous background parameters are the same as in the previous section. For author's best knowledge, to date, the crosstalk problem between μ_s and g has not been considered in the literature for the DOT. Hence, we are interested to study the crosstalk effect of the scattering coefficient on the anisotropy factor reconstruction, and vice-versa. For this purpose, two different test cases are considered. In the first one, the inclusion A varies only in scattering ($\mu_s^* = 4 \text{ mm}^{-1}$) while the inclusion B represents a heterogeneity in anisotropy factor ($g^* = 0.85$). In the second test, the inclusion A is an anisotropy

heterogeneity only ($g^* = 0.85$) and the inclusion B varies only in scattering ($\mu_s^* = 4 \text{ mm}^{-1}$). This last is made in order to show the influence of the inclusion location with respect to the source on the reconstruction results. The reconstructed images are displayed in Fig. 5 and the relative errors are listed in Table 4.

[Figure 5 about here.]

From Table 4, the reconstruction quality of the inclusion A is relatively more accurate ($\varepsilon_{\mu_s}^{IncA} = 17.20\%$ and $\varepsilon_g^{IncA} = 3.00\%$) than that of the inclusion B ($\varepsilon_{\mu_s}^{IncB} = 19.00\%$ and $\varepsilon_g^{IncB} = 4.35\%$) in scattering and anisotropy factor. This is mainly due to the location effect, since the inclusion A is nearer to the source and placed in higher sensitivity area than the inclusion B. Therefore, the inclusion A transmits more rich informations to detectors and makes the inversion more accurate. Unlike the cases where μ_a and μ_s were retrieved simultaneously, the crosstalk issue between the scattering and the anisotropy factor is clearly pronounced in both optical images. This is because μ_s and g have both significative and important sensitivities on the emerging intensities. Comparing the two test cases, it can be observed that the crosstalk impact depends on the inclusion location with respect to the source. The crosstalk μ_s error has decreased from $\varepsilon_{\mu_s}^{Crosstalk} = 9.66\%$ to $\varepsilon_{\mu_s}^{Crosstalk} = 8.62\%$ when the g heterogeneity was exchanged from inclusion A to B. Similarly, the crosstalk g error has decreased from $\varepsilon_g^{Crosstalk} = 1.03\%$ to $\varepsilon_g^{Crosstalk} = 0.49\%$ when the scattering inclusion passed from position A to B. Thus, we can deduce that the crosstalk is more pronounced as the responsible inclusion in the other parameter is nearer to the source (inclusion A). Note that, the μ_s inclusion B (case 2) has a weak crosstalk effect on the g image (Fig. 5d) whereas when this inclusion is a g heterogeneity (case 1), the crosstalk in the μ_s image is much more remarkable (Fig. 5a). This can be attributed to the different sensitivities for μ_s and g on the reflectance. It has been shown in [43] that the reflectance of the RTE model is much more sensitive to a variation in the anisotropy factor than the scattering coefficient. That explains the better estimation quality of g for all cases (Tab. 4) and further the strong crosstalk in the reconstructed μ_s images. Another interesting remark can be observed when μ_s and g are reconstruct simultaneously. As shown in Figs. 5(a,b,c,d), the low inclusion in anisotropy factor appears as a high scattering inclusion in the μ_s images. Also, the low scattering inclusion is reconstructed as a high inclusion in anisotropy factor. Hence, we can deduce that the two parameters μ_s and g doesn't have the same behavior and effect on the backscattered light. These parameters present an opposite sensitivities on the reflectance [6, 43]. In other words, an increase in the reflectance could be caused by either a decrease in g factor or an increase in μ_s coefficient, and vice-versa. In case 1 for example, the increase of the reflectance due to the inclusion B (low anisotropy factor $g^* = 0.85$) has resulted in low heterogeneity in g (Fig. 5b) and highly scattering inclusion (crosstalk) (Fig. 5a). The reconstructed results are in agreement with the previous sensitivity studies and confirm the non-uniqueness nature of the problem when several combinations of μ_s and g can lead to identical

boundary data.

[Table 4 about here.]

4.5.2 Without crosstalk problem

For computational time purpose, the simulated reconstructions for all the previous cases were performed on a relatively small phantom size. In this subsection, we consider a $2\text{ cm} \times 2\text{ cm}$ domain which contains one circular inclusion in order to test our algorithm for larger domain. The source-detectors configuration is the same as in the previous subsection in order to get closer to a realistic experimental setup based on the reflectance geometry. The synthetic data are generated on a finer triangular mesh of 21248 elements using the exact optical properties while the inversion is performed on a coarser mesh of 5312 triangular elements. The data are corrupted by 3% of noise level which is inevitable in the practical applications. The inclusion located at position $x = 5\text{ mm}$, $y = 7\text{ mm}$ is assigned the exact values $\mu_s^* = 6\text{ mm}^{-1}$ and $g^* = 0.85$ while $\mu_a = 0.05\text{ mm}^{-1}$ is uniformly distributed inside the medium. The reconstructed results are depicted in Fig. 6.

[Figure 6 about here.]

As shown in Figure 6, the inclusion is accurately identified in the exact location for both optical properties. It is also seen that the reconstructed profiles of μ_s and g along the cross-section on $y = 7\text{ mm}$ are spatially well fitted with the exact solution (Figs. 6 (c,d)). Furthermore, the values of optical properties are accurately retrieved even for noisy data. Also, the background is well recovered for μ_s and g images. These results prove that the algorithm is robust for realistic domain size and can provide a good estimation quality for both μ_s and g in this situation. However, if the exact values μ_s^* and g^* of the inclusion are both either high or low with respect to the background values, the quality reconstruction will be relatively worse. This behavior is due to the opposite sensitivities of the two parameters on the boundary data, as deduced previously. The residual errors induced by this inclusion will be compensated yielding to small objective function level and therefore to image inaccuracy. Hence, a suitable regularization technique for the proposed algorithm is needed in order to better separate and estimate simultaneously the optical properties μ_s and g .

5 Summary

Reconstructions of spatial distributions of the optical properties for different cases were presented. The radiative transfer equation was used as forward model in frequency-domain and solved accurately by the MFVM. For the inversion, the gradient of the objective function with respect to μ_a , μ_s and g was computed fastly and efficiently by using the adjoint

method. The simultaneous reconstructions of μ_a and μ_s were achieved with reasonable accuracy for lowly and even for highly absorbing media. It has also been pointed out that the estimation and crosstalk errors depend on both the inclusion location and contrast level. The main contribution of this study is the reconstruction of the spatial distribution of the anisotropy factor g . This estimation of g was possible by using forward and adjoint models based on the RTE. Consequently, this work might open the possibility to image g in tissues as an additional contrast agent for DOT. Also, we have shown the capability of the proposed algorithm to reconstruct simultaneously μ_s and g even for large domain with noisy data. The crosstalk between the two parameters has been considered and clearly observed in both optical images. We deduced that μ_s and g have an opposite sensitivities and effects on the reflectance. Therefore, a suitable regularization technique will be implemented as a next step, in order to reduce the strong crosstalk issue between μ_s and g . This work was a necessary preliminary study before extending the present algorithm to 3D reconstructions.

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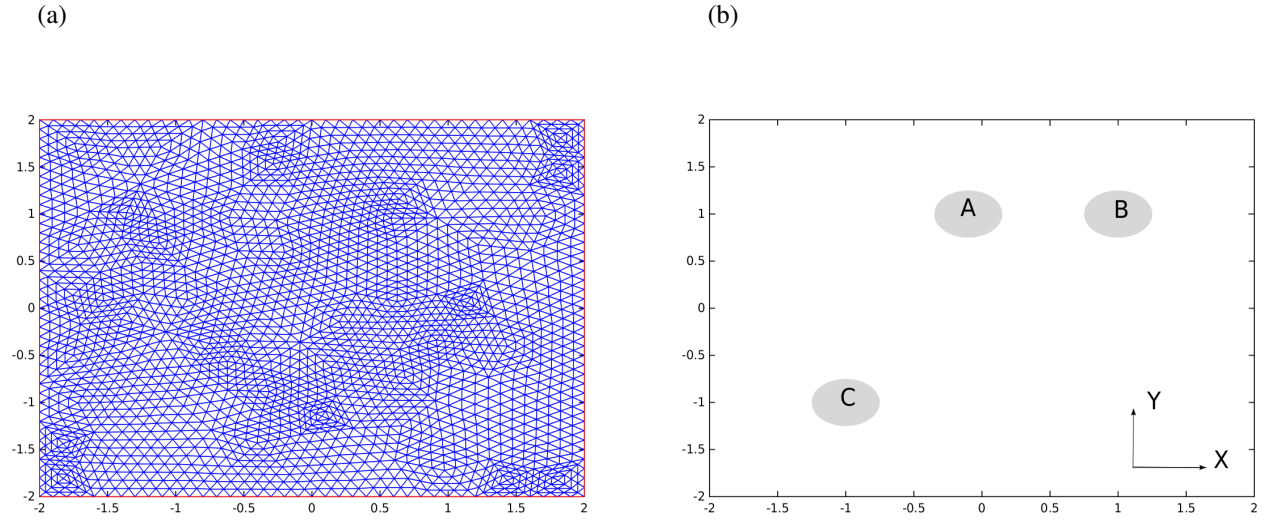


Figure 1: (a) Two-dimensional ($4 \times 4 \text{ mm}^2$) triangular mesh (b) containing three circular inclusions A, B and C.

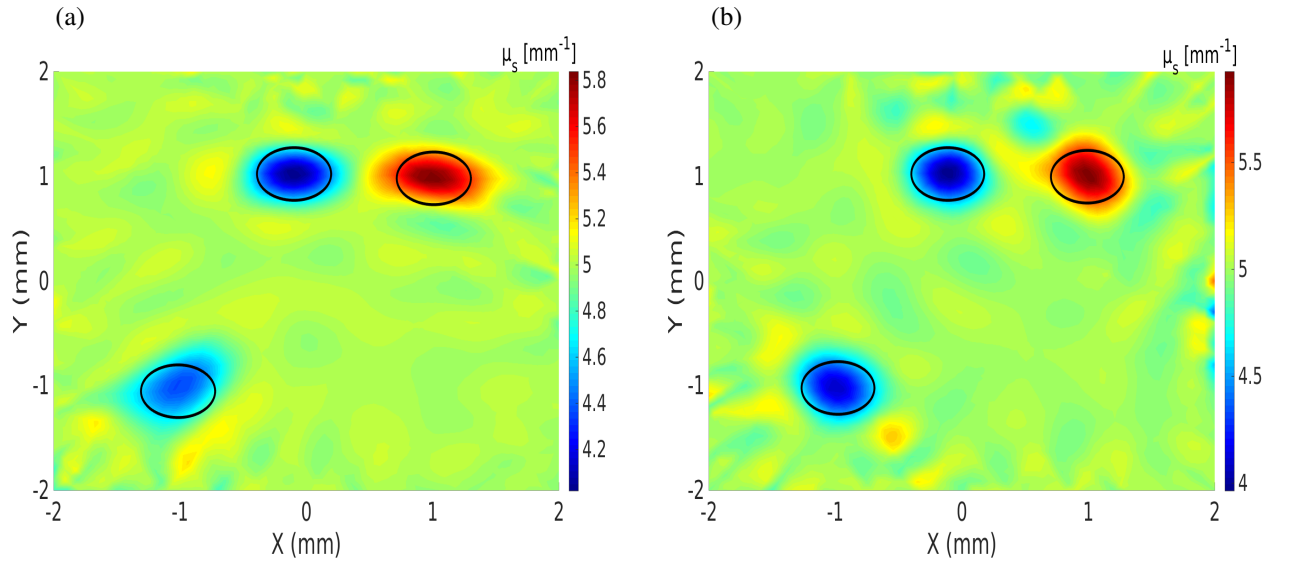


Figure 2: Reconstructions of the scattering coefficient μ_s for two different source numbers. (a) One source is placed at the mid-center of the north surface (b) Four sources are used to probe the mid-center of each surface of the phantom. The solid circles indicate the exact inclusion locations.

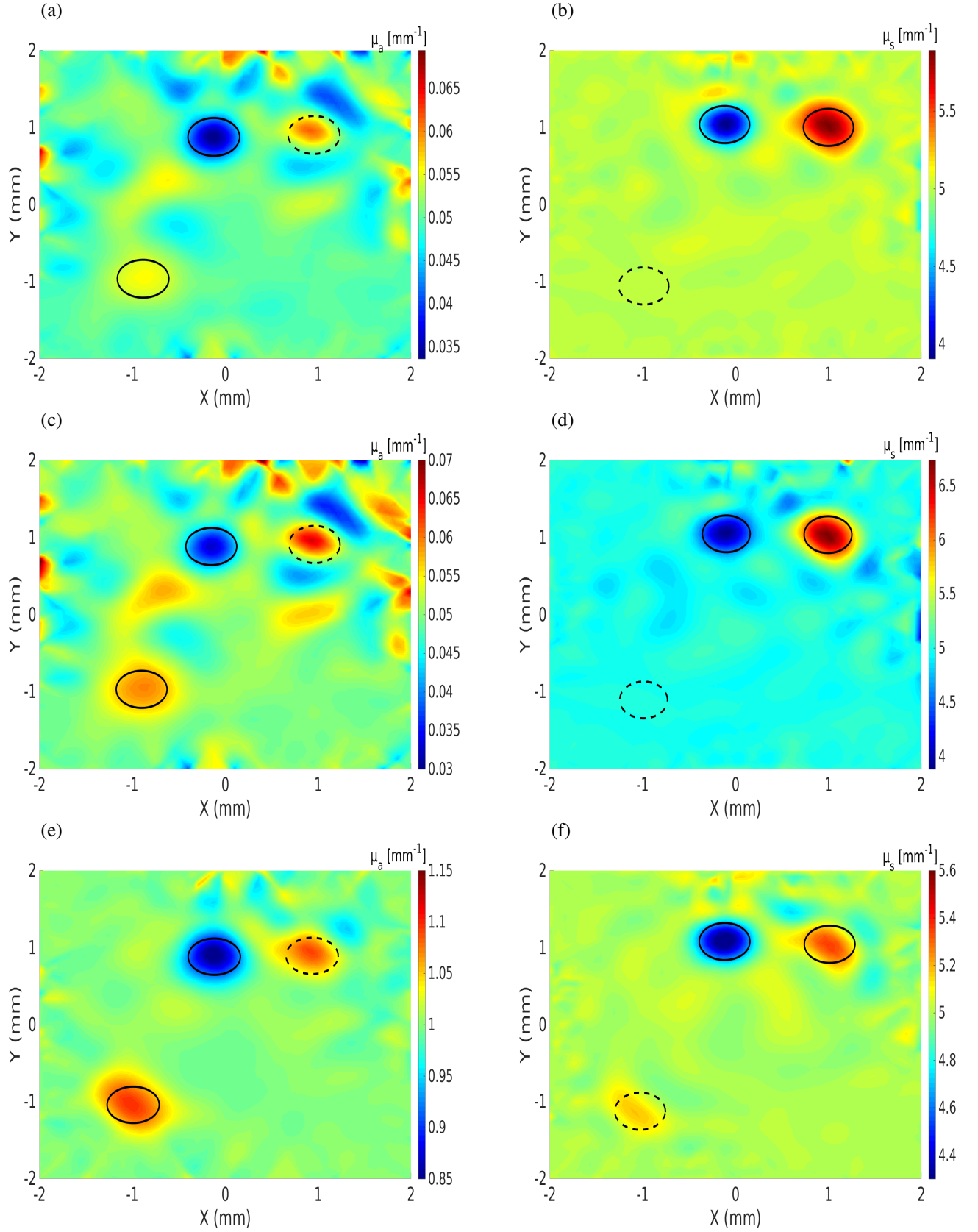


Figure 3: Simultaneous reconstructions of the absorption μ_a and the scattering μ_s coefficients. Left column : Reconstructed μ_a images. Right column : Reconstructed μ_s images. Top row : Test case 1. Middle row : Test case 2. Bottom row : Test case 3. The solide circles indicate the exact positions while the dashed circles depict the crosstalk zones. We started the minimization using the homogeneous background optical properties.

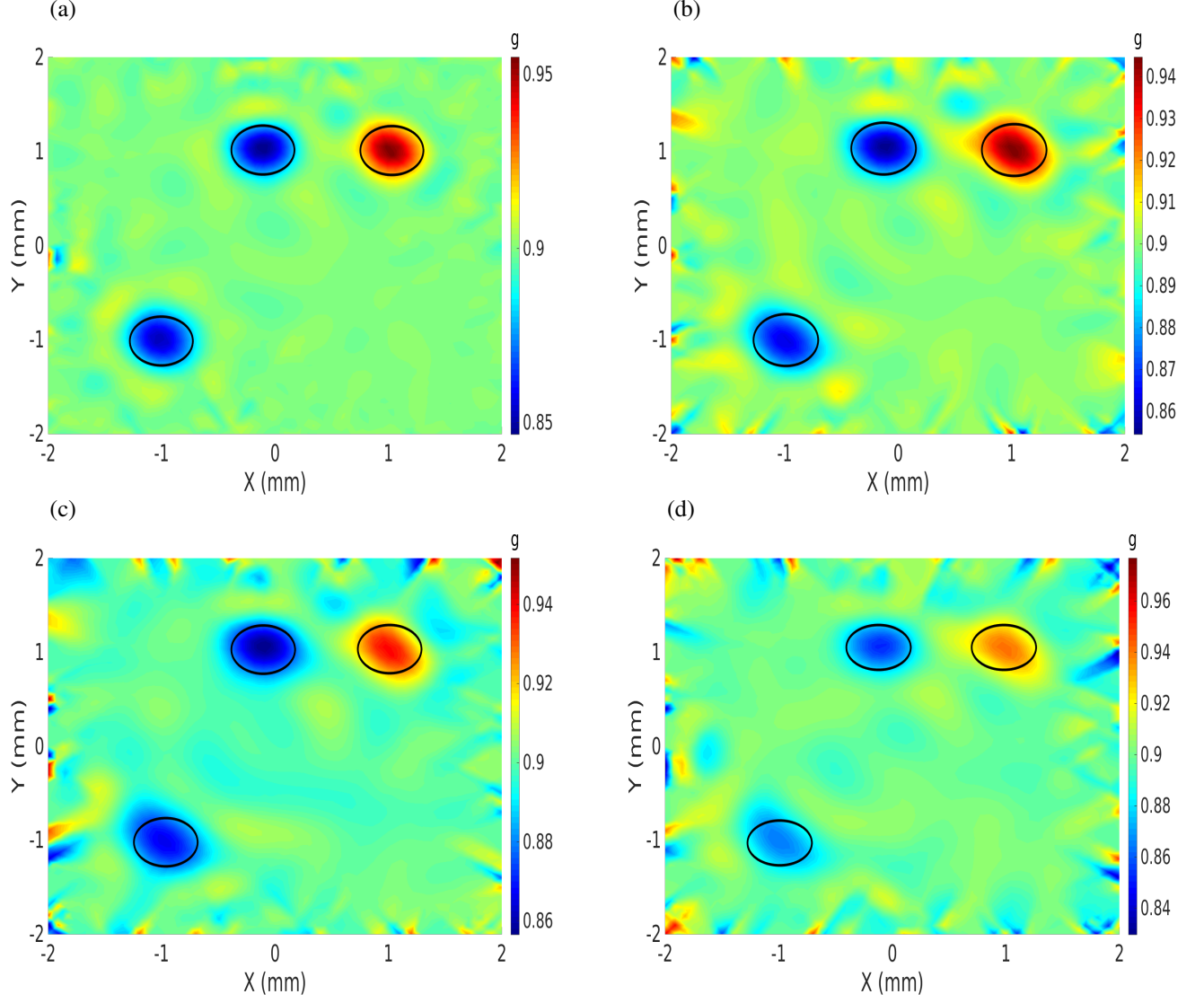


Figure 4: Reconstructions of the anisotropy factor g for four different noise levels σ_m on the synthetic data. The original phantom medium to reconstruct was shown in figure 1. (a) reconstructed g image with noiseless data (b) reconstructed g image, for $\sigma_m = 3\%$ (c) reconstructed g image, for $\sigma_m = 6\%$ (d) reconstructed g image, for $\sigma_m = 10\%$. The solide circles indicate the exact inclusion locations.

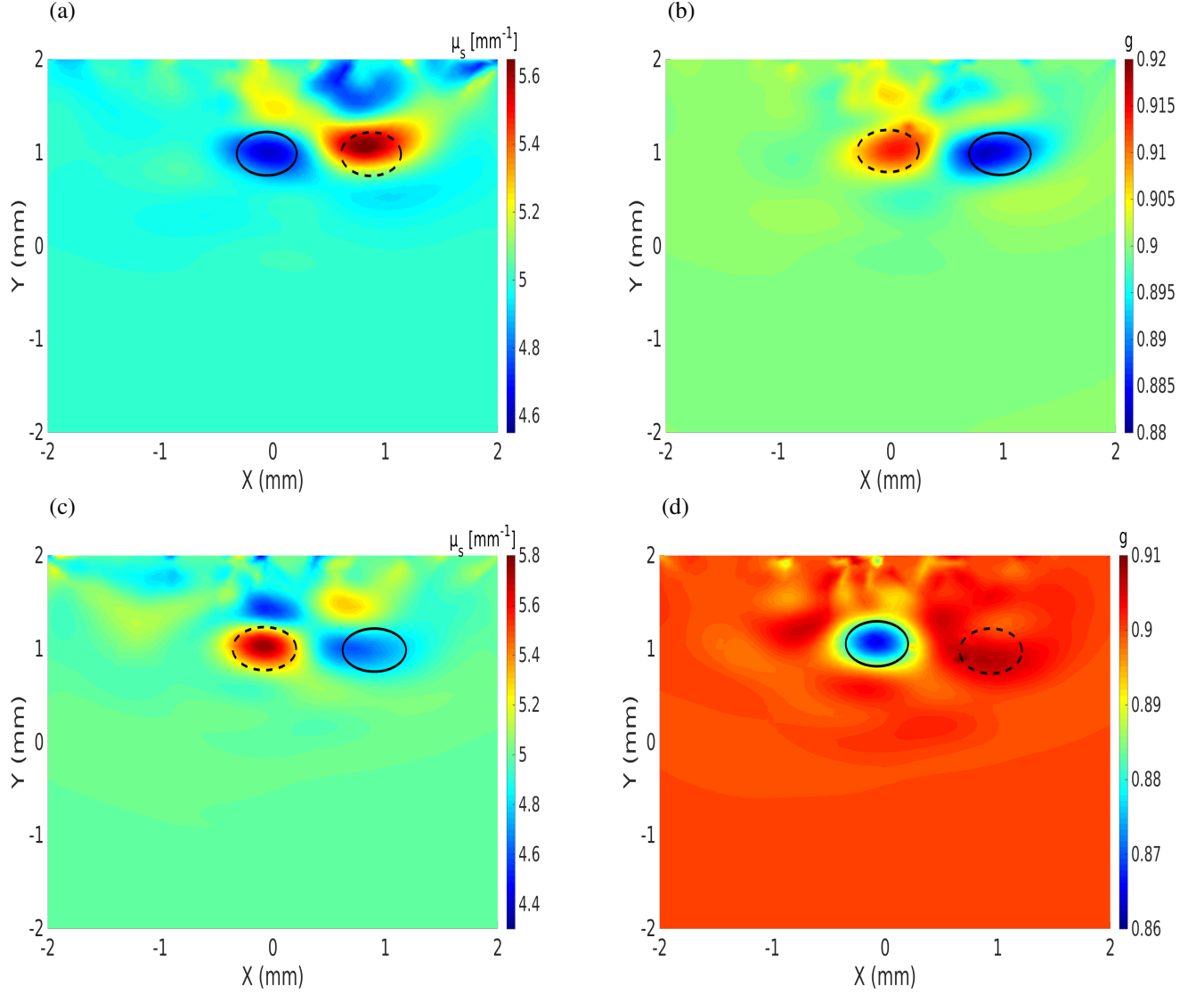


Figure 5: Simultaneous reconstructions of the scattering coefficient μ_s and the anisotropy factor g . Left column : Reconstructed μ_s images. Right column : Reconstructed g images. Top row : Test case 1, inclusion A in scattering coefficient and inclusion B in anisotropy factor. Bottom row : Test case 2, inclusion A in anisotropy factor while inclusion B in scattering coefficient. The solid circles indicate the exact positions while the dashed circles depict the crosstalk zones.

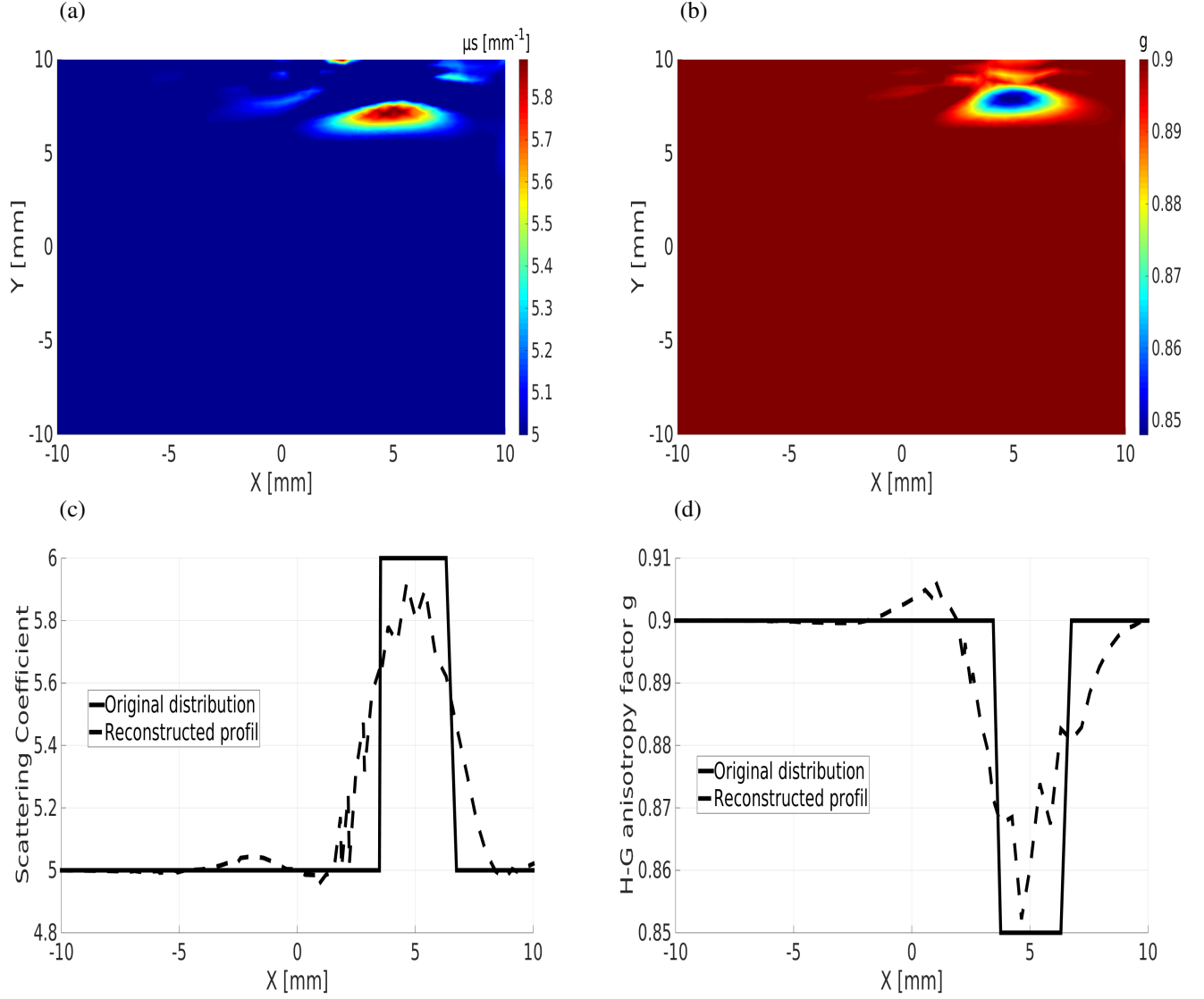


Figure 6: Simultaneous reconstructions of the scattering coefficient μ_s and the anisotropy factor g for the $2 \text{ cm} \times 2 \text{ cm}$ domain. (a) : Reconstructed μ_s image. (b) : Reconstructed g image. (c) : μ_s Cross-section. (d) : g Cross-section. We started the minimization using the homogeneous background optical properties : $\mu_a = 0.05 \text{ mm}^{-1}$, $\mu_s = 5 \text{ mm}^{-1}$ and $g = 0.9$.

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		Inclusions			
		Background	A	B	C
Test case 1	$\mu_a(\text{mm}^{-1})$	0.05	0.04	0.05	0.06
	$\mu_s(\text{mm}^{-1})$	5	4	6	5
Test case 2	$\mu_a(\text{mm}^{-1})$	0.05	0.04	0.05	0.07
	$\mu_s(\text{mm}^{-1})$	5	4	7	5
Test case 3	$\mu_a(\text{mm}^{-1})$	1	0.8	1	1.2
	$\mu_s(\text{mm}^{-1})$	5	4	6	5

Table 1: The exact absorption and scattering coefficients of the 3 test mediums

	μ_a				μ_s			
	$\epsilon_{\mu_a}^{background}$	$\epsilon_{\mu_a}^{IncA}$	$\epsilon_{\mu_a}^{IncC}$	$\epsilon_{\mu_a}^{Crosstalk}$	$\epsilon_{\mu_s}^{background}$	$\epsilon_{\mu_s}^{IncA}$	$\epsilon_{\mu_s}^{IncB}$	$\epsilon_{\mu_s}^{Crosstalk}$
Test medium 1	3.57%	10.42%	9.16%	12.29%	0.75%	6.88%	5.46%	0.17%
Test medium 2	6.11%	10.95%	17.7%	24.49%	1.36%	7.35%	9.70%	0.27%
Test medium 3	0.94%	11.62%	9.94%	5.769%	0.60%	12.1%	12.6%	1.85%

Table 2: The relative estimation errors of background, inclusions and crosstalk for the absorption and scattering coefficients.

	Errors			
	$\sigma_m = 0\%$	$\sigma_m = 3\%$	$\sigma_m = 6\%$	$\sigma_m = 10\%$
Relative estimation error ϵ_g (%)	0.29	0.41	0.53	0.72
Number of iterations k	57	22	17	15

Table 3: The relative estimation errors and the iterations numbers of the reconstruction algorithm for the 4 different noise levels.

	μ_s				g			
	$\varepsilon_{\mu_s}^{background}$	$\varepsilon_{\mu_s}^{IncA}$	$\varepsilon_{\mu_s}^{IncB}$	$\varepsilon_{\mu_s}^{Crosstalk}$	$\varepsilon_g^{background}$	ε_g^{IncA}	ε_g^{IncB}	$\varepsilon_g^{Crosstalk}$
Test case 1	0.62%	17.20%	-	8.62%	0.08%	-	4.35%	1.03%
Test case 2	0.67%	-	19%	9.66%	0.1%	3%	-	0.49%

Table 4: The relative estimation errors of background, inclusions and crosstalk for the scattering coefficient μ_s and the anisotropy factor g .