Forward Problem Study of an Effective Medium Model for Ultrasound Blood Characterization

Emilie Franceschini, Member, IEEE, Bloen Metzger, and Guy Cloutier, Senior Member, IEEE

Abstract—The structure factor model (SFM) is a scattering model developed to simulate the backscattering coefficient (BSC) of aggregated red blood cells (RBCs). However, the SFM can hardly be implemented to estimate the structural aggregate parameters in the framework of an inverse problem formulation. A scattering model called the effective medium theory combined with the SFM (EMTSFM) is thus proposed to approximate the SFM. The EMTSFM assumes that aggregates of RBCs can be treated as individual homogeneous scatterers, which have effective properties determined by the acoustical characteristics and concentration of RBCs within aggregates. The EMTSFM parameterizes the BSC by three indices: the aggregate radius, the concentration of RBCs within aggregates (the aggregate compactness), and the systemic hematocrit. The goodness of fit of the EMTSFM approximation in comparison with the SFM was then examined. Based on a 2-D study, the EMTSFM was found to approximate the SFM with relative errors less than 30% for a product of the wavenumber times the mean aggregate radius $kr_{\rm ag} \leq 1.32$. The main contribution of this work is the parameterization of the BSC with the RBC aggregate compactness, which is of relevance in clinical hemorheology because it reflects the binding energy between RBCs.

I. INTRODUCTION

ULTRASONIC tissue characterization techniques using the RF backscattered signals have received broad interest for the past 25 years. One approach is to use the magnitude and frequency dependence of backscatter echoes to quantify the tissue structures such as the size, acoustic impedance, and concentration of the scatterers. This approach has been successfully used for the characterization of the eye [1], the prostate [2], apoptotic cells [3], and the breast [4]. Blood has also been studied by employing this technique [5], [6]. In the field of ultrasonic blood characterization, the objective is to assess the level

B. Metzger is with the Institut Universitaire des Systèmes Thermiques Industriels (IUSTI), Centre National de la Recherche Scientifique (CNRS), Marseille, France.

G. Cloutier is with the Laboratory of Biorheology and Medical Ultrasonics, University of Montreal Hospital Research Centre (CRCHUM), Montreal, Quebec, Canada, and with the Department of Radiology, Radio-Oncology, and Nuclear Medicine, and the Institute of Biomedical Engineering, University of Montreal, Montreal, Quebec, Canada.

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of red blood cell (RBC) aggregation, which is a surrogate marker of inflammation [7]. It is well known that when RBCs are under low shear rates ($<10 \text{ s}^{-1}$), they interact strongly with each other and form complex rouleaux, or 3-D structures. When the shear rate increases, these rouleaux or structures disaggregate. The aggregation phenomenon in human blood is normal, however hyperaggregation, an abnormal increase of RBC aggregation, is a pathological state associated with several circulatory diseases such as deep venous thrombosis, atherosclerosis, and diabetes mellitus. Ultrasonic blood characterization using the ultrasound backscatter technique has the potential to provide a method for the noninvasive estimation of the RBC aggregate size. This quantification would help to elucidate the role of RBC aggregation in the etiology of such diseases.

Ultrasound backscattering by blood is mainly due to RBCs that constitute the vast majority (97%) of the blood cellular content. Blood can thus be described as a biphasic fluid composed of RBCs immersed in plasma. Because RBCs are acoustically considered to be weak scatterers (impedance contrast between RBCs and plasma being around 13%), multiple scattering can be neglected. However, for such tissue, it is not straightforward to develop a theoretical scattering model because of the high density of RBCs (their volume fraction or hematocrit varies between 30 and 50%) and their ability to form aggregates. Theoretical efforts have been made to take into account the high density of RBCs [5], [8]–[10]. In the Rayleigh scattering regime (i.e., for a product of the wavenumber times the scatterer radius $ka \ll 1$), Twersky [9] proposed an expression of the backscattered intensity in terms of the single-particle backscattering cross section, number density of particles and packing factor. The packing factor is dependent on the hematocrit but independent of the frequency. This model succeeded in explaining the nonlinear relationship between the backscatter amplitude and hematocrit for non-aggregating RBCs [11] but failed to predict the magnitude and frequency dependence of backscatter echoes observed in *in vitro* experiments when considering aggregating RBCs. That is why Savery and Cloutier [12] proposed generalization of the packing factor theory for aggregating RBCs at a low hematocrit by introducing the frequency-dependent structure factor, named the structure factor model (SFM). This model was later generalized to a normal hematocrit of 40% [13]. The SFM sums the contributions from individual RBCs and models the RBC interaction by a statistical mechanics structure

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E. Franceschini is with the Laboratoire de Mécanique et d'Acoustique (LMA), Centre National de la Recherche Scientifique (CNRS) UPR 7051, Marseille, France (e-mail: franceschini@lma.cnrs-mrs.fr).

TABLE I. ACOUSTICAL PROPERTIES	of Blood Fo	und in [11	AND	[21]
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	Density $\rho \; (\text{kg} \cdot \text{m}^{-3})$	Compressibility $\kappa (Pa^{-1})$	Impedance Z (MRayl)
RBC*	1092	3.41×10^{-10}	1.766
Plasma	1021	4.09×10^{-10}	1.580
*RBC = red blood	cells.		

factor, which is, by definition, the Fourier transform of the spatial distribution of RBCs [12]-[14]. Note that the lowfrequency limit of the structure factor is, by definition, the packing factor used under Rayleigh conditions, and that the structure factor cannot analytically be calculated contrary to the packing factor [9]. The SFM was largely used to perform simulation studies on RBC aggregation [12], [13], [15]–[17]. Simulations enabled prediction of the frequency dependence of the backscattering coefficient (BSC) from various RBC spatial distributions and thus helped the interpretation of experimental observations. However, the SFM cannot directly be used to estimate the RBC aggregate size in the framework of an inverse problem approach because of the intensive computational time required to assess the structure factor by realizing distributions of aggregating RBCs with simulations. Note that the SFM should not be confused with the structure factor size estimator (SFSE) recently proposed by Yu et al. [18], [19]. The SFSE approximates the SFM for practical assessments of RBC structural features (i.e., in an inverse problem formulation).

Yu et al. [18], [19] developed the SFSE scattering theory that approximates the SFM by using a second-order Taylor expansion of the structure factor. The SFSE is thus not as accurate as the SFM. The SFSE parameterizes the BSC by two structure indices: the packing factor and the mean aggregate diameter, assumed to be isotropic. However, experiments with pig blood in controlled flow devices [18], [19] and 3-D numerical simulations of isotropic aggregates [16] showed that the two indices are correlated and follow a quadratic relationship under the assumption of isotropic monodisperse aggregates, thus reducing the BSC parameterization to one structural index. Moreover, the SFSE model did not produce good fits to the simulated BSCs computed with the SFM [16, Fig. 4]. Thus, the SFSE model is not sufficient to approximate the SFM accurately; even if the relation between the simulated aggregate sizes and its SFSE parameterization followed a proportional relationship [16]. The goal of this paper is to propose a new scattering model that better approximates the SFM for structural RBC aggregate characterization.

The scattering theory we propose is based on an effective medium theory (EMT) combined with the SFM, labeled EMTSFM. The EMT was initially developed by Kuster and Toksoz [20] in the field of geophysics. Here, the EMT assumes that aggregates of RBCs can be treated as individual homogeneous scatterers, which have effective properties determined by the concentration of RBCs within aggregates and acoustical properties of blood constituents. The approximation of RBC aggregates as ho-

mogeneous effective particles is combined with the SFM to consider the concentrated blood medium. The effective particle interactions were thus modeled by a structure factor, as in [12] and [13]. The EMTSFM parameterizes the BSC by three indices: the aggregate radius, the internal hematocrit (i.e., the concentration of RBCs within aggregates, also called the aggregate compactness) and the systemic hematocrit. Note that the new EMTSFM model would allow characterizing, for the first time in the quantitative ultrasound field, the compactness of RBC aggregates (or of any other cellular structures). In the field of clinical hemorheology, assessing the compactness of RBC aggregates is of high clinical importance because it is related to the binding energy between cells. Normal RBC aggregates form rouleaux-type structures, whereas pathologies associated with stronger binding energies result in clumps of RBCs (close to a spherical isotropic packing).

The purpose of this study was to compare the EMTSFM and SFM in the framework of a forward problem study, i.e., determining the BSC from known distributions of RBCs with known acoustical parameters using both EMTSFM and SFM. The goodness of fit of the EMTSFM approximation in comparison with the SFM was examined as a function of the frequency and of the structural aggregate parameters (i.e., the aggregate size and the internal hematocrit). Because of the computational load required to generate 3-D RBC distributions with various internal hematocrits with the SFM, 2-D models were used for the BSC computation. Although limiting, this choice allowed us to simulate randomly various internal hematocrits for a given range of systemic hematocrits, and to isolate the effects of aggregate size and internal hematocrit on the BSC.

II. ULTRASOUND BACKSCATTERING THEORY

In the following, it is assumed that the incident wavelength λ is large compared with the RBC size. Consequently, the RBC shape could be approximated by a simple geometry having an equivalent volume of an RBC in 3-D or having an equivalent surface in 2-D [21]. In the present 2-D study, RBCs were modeled as parallel infinite cylinders of radius *a*, that have small contrast in acoustical properties relative to the plasma (see Table I). Two scattering models of RBC aggregation are presented in this section: the SFM and the new EMTSFM.

A. The Structure Factor Model (SFM)

The SFM of ultrasound backscattering by blood consists of summing contributions from individual RBCs and modeling the RBC interaction by a statistical mechanics structure factor [12], [13], [15]. By considering a collection of N identical and weakly scattering RBCs, the BSC expression can be written as

$$BSC_{SFM}(-2\mathbf{k}) = m\sigma_b(-2\mathbf{k})S(-2\mathbf{k}), \qquad (1)$$

where **k** is the incident wave vector and *m* is the number density of RBCs that is related to the systemic hematocrit ϕ as $m = \phi/A_p$ (where A_p is the RBC area). The backscattering cross section σ_b of a single weak RBC can be written as the product of the backscattering cross section of a single RBC in the Rayleigh limit [9, Eq. (16)] and the backscatter form factor *F* as

$$\sigma_b(-2\mathbf{k}) = \frac{k^3 A_p^2 \gamma_z^2}{2\pi} F(-2k, a) = \frac{k^3 A_p^2 \gamma_z^2}{2\pi} \Big(\frac{J_1(2ka)}{ka}\Big)^2, \ (2)$$

where J_1 is the first-order Bessel function of the first kind and $\gamma_z = (Z_{\text{RBC}} - Z_{\text{plasma}})/Z_{\text{plasma}}$ is the fractional variation of impedance between the RBC and its suspending medium (i.e., the plasma). The form factor F serves as a useful descriptor of a scatterer with a simple shape [21], [22] and characterizes here the cylindrical shape and size of the 2-D RBC scatterer. The demonstration of the cylindrical form factor expression in 2-D is given in the Appendix. The function S in (1) is the structure factor representing the spatial positioning of RBCs; it is defined by

$$S(-2\mathbf{k}) = E\left[\frac{1}{N}\left|\sum_{i=1}^{N} e^{-i2\mathbf{k}\mathbf{r}_{i}}\right|^{2}\right],$$
(3)

where E is the expected value of a random variable and \mathbf{r}_i is the position vector defining the center of the *i*th RBC in space. Note that the aggregation phenomenon only affects the structure factor because RBC properties (i.e., σ_b) and the systemic hematocrit are expected to remain constant in the modeled region of interest.

In the case of disaggregated RBCs, the low-frequency limit of the structure factor is a constant with this model $[S(k) \rightarrow S(0) = W]$, which is the packing factor. The BSC expression can thus be simplified as

$$BSC_{LF}(-2\mathbf{k}) = m W \sigma_b(-2\mathbf{k}). \tag{4}$$

The most often used packing factor expression is based on the Percus-Yevick pair-correlation function for identical, hard, and radially symmetrical particles. The Percus-Yevick packing factor $W_{\rm PY}$ was first applied to blood by Shung *et al.*, and it is related to the systemic hematocrit in the 2-D space as [11]

$$W_{\rm PY} = \frac{(1-\phi)^3}{1+\phi}.$$
 (5)

The packing factor reflects the positional correlation among particles: W = 1 for completely random distribu-



Fig. 1. Schematic representation of aggregates treated as individual scatterers. The aggregates of red blood cells in blood (left side) are assumed to be homogeneous particles (right side) with effective properties that depend on the internal hematocrit and on the density and compressibility of the red blood cells within them.

tions and W tends toward zero as the particle number and the spatial correlation between particles increase.

B. The Proposed Model: An Effective Medium Theory Combined With the SFM (EMTSFM)

As a first approximation, we assume that all the RBCs are aggregated in blood, that the aggregates are identical and isotropic and that the RBCs within the aggregates are evenly distributed. The EMTSFM assumes that aggregates of RBCs can be treated as individual homogeneous scatterers as shown in Fig. 1. Each aggregate is thus approximated by an effective single particle (i.e., in this 2-D study, an effective single cylinder) having a radius $r_{\rm ag}$. The density $\rho_{\rm ag}$ and compressibility $\kappa_{\rm ag}$ of the new effective particle are determined by considering the EMT [20]. This means that ρ_{ag} and κ_{ag} are derived from the acoustical properties of the two fluids that constitute the aggregates (i.e., ρ_1 , ρ_2 , κ_1 , and κ_2 , where 1 indicates properties of RBCs and 2 those of plasma) and from the internal concentration of RBCs within the aggregates, defined as the internal hematocrit ϕ_i , as follows:

$$\rho_{\rm ag} = \phi_i \rho_1 + (1 - \phi_i) \rho_2$$

$$\frac{1}{\kappa_{\rm ag}} = \frac{\phi_i}{\kappa_1} + \frac{1 - \phi_i}{\kappa_2}.$$
(6)

The acoustic interaction of RBCs within aggregates is therefore taken into account with the EMT, similarly to the SFM, where it is considered by the structure factor that models the individual position of RBCs and their acoustical interaction, whether they are within an aggregate or not. The main advantage of the EMTSFM is the consideration of the compactness of aggregates with the parameter ϕ_i .

The BSC from blood is then obtained by summing contributions from individual effective particles of radius $r_{\rm ag}$ and modeling the effective particle interaction by a statistical mechanics structure factor, $S_{\rm ag}$. The equivalent BSC expression is thus

$$BSC_{eq}(-2\mathbf{k}) = m_{ag}\sigma_{ag}(-2\mathbf{k})S_{ag}(-2\mathbf{k}), \qquad (7)$$

where $m_{\rm ag}$ is the number density of aggregates, which is related to the effective area fraction of aggregates, $\phi_{\rm ag}$. The effective area fraction of aggregates is equal to the area fraction of RBCs in blood, ϕ , divided by the internal hematocrit ϕ_i : $\phi_{ag} = \phi/\phi_i$. The backscattering cross section, σ_{ag} , of an effective single cylinder was calculated using the fluid-filled cylinder model developed by Doolittle and Uberall [23]. That model provides an exact solution for the backscattering of sound by a single fluid cylinder, not necessarily small compared with the wavelength, in a surrounding fluid medium (i.e., the plasma). In the proposed forward problem study, the radius $r_{\rm ag},$ density $\rho_{\rm ag},$ and compressibility κ_{ag} of the aggregates, and the density, ρ_2 , and compressibility, κ_2 , of the plasma are known a priori (see Table I) such that the backscattered pressure of the effective fluid cylinder could be computed as a function of frequency. The structure factor $S_{\rm ag}$ was calculated for a collection of $N_{\rm ag}$ identical and disaggregated particles (mimicking RBC aggregates) of radius $r_{\rm ag}$, randomly distributed as follows:

$$S_{\rm ag}(-2\mathbf{k}) = E \left[\frac{1}{N_{\rm ag}} \left| \sum_{i=1}^{N_{\rm ag}} e^{-i2\mathbf{k}\mathbf{r}'_i} \right|^2 \right], \tag{8}$$

where \mathbf{r}'_i are the position vectors defining the center of the *i*th effective particle (or aggregate) in space. Note that in the research field of optics [24], the applicability of the modeling approach based on the structure factor (called the interference approximation) is limited to a product $kR \leq 3.5$ (or $kR \leq 1.5$, respectively) for a relative refractive index of the particles equal to 1.19 (or 1.8), where R is the radius of isotropic identical particles. Because the relative acoustic impedance of RBCs, z_1/z_2 , is around 1.12 (see Table I), we are thus expecting the same limitation for the EMTSFM (i.e., a validity of this modeling approach restricted to $kr_{ag} \leq 3.5$).

In the low-frequency limit, for aggregate sizes which are small compared with the wavelength, the structure factor $S_{\rm ag}$ can be well approximated by the Percus-Yevick packing factor of (5) for effective particles. The equivalent BSC expression is thus simplified as

$$BSC_{eq_{LF}}(-2\mathbf{k}) = m_{ag} \frac{(1-\phi_{ag})^3}{1+\phi_{ag}} \sigma_{ag}(-2\mathbf{k}), \qquad (9)$$

where σ_{ag} is calculated using the exact model of fluid-filled cylinders [23], such that, in comparison with (7), the only approximation is the structure factor S_{ag} .

III. SIMULATION METHODS

The computation of the BSC_{SFM} and BSC_{eq} requires the knowledge of the structure factors S and S_{ag} , as described in (1) and (7). Because the structure factor S (or equivalently S_{ag}) is, by definition, a statistical quantity, an average of all structure factors obtained from several particle distributions can give an estimated value of S (or equivalently S_a). Note that 2-D models were used because of the computational time required to generate 3-D RBC distributions with the SFM. Although limiting, this choice allowed us to simulate randomly varied internal hematocrits for a given range of systemic hematocrits.

A. Computation of Particle Distributions

1) Distribution of RBCs for the Computation of S With the SFM: We describe here how random distributions for non-aggregating and aggregating RBCs were computed within the simulated surface area L^2 of $600^2 \,\mu\text{m}^2$. The RBC particle radius a was set to 2.75 μ m for all simulations. In the case of disaggregated RBCs, particles were randomly distributed (using a random number generator) with non-overlapping positions to give the desired area fraction of RBCs in blood (i.e., the systemic hematocrit ϕ). In the case of aggregated RBCs, we first specified the aggregate radius and then aggregates were randomly distributed with non-overlapping positions to give the desired area fraction of aggregates ϕ_{ag} . RBC distributions within aggregates were then generated by considering the number of RBCs within aggregates n_i and the area fraction of RBCs within aggregates ϕ_i . It is important to emphasize that random particle distributions could be easily generated using a random number generator up to an area fraction of approximately 0.5. In our study, the area fraction of RBCs within aggregates ϕ_i could be enlarged, up to a maximum value $\phi_{i_{\rm max}}$ fixed at 0.6. The procedure we chose to reach such a high number of RBCs within aggregates is described in the following.

We first randomly distributed n_i RBCs in a total area that was larger than the aggregate area, such that the initial area fraction of RBCs within aggregates was lower than 0.5. Then, to increase this area fraction, the RBCs were submitted to an external force oriented toward the center of mass of the RBC spatial distribution. RBCs moved toward each other while a repulsive force prevented them from overlapping. These forces were maintained until the area fraction of RBCs inside the considered aggregate reached the desired area fraction ϕ_i . This distribution procedure was repeated for each aggregate, such that the distribution of RBCs within each aggregate was different. Note that this procedure is not time consuming if a small number of RBCs is used: herein the maximum number of RBCs within aggregates was 38 for the maximum aggregate radius of $7.95a = 21.86 \ \mu\text{m}$. The maximum value of aggregate area fraction $\phi_{\mathrm{ag}_{\mathrm{max}}}$ was thus fixed to 0.5 and as a consequence, the maximum value of the systemic hemato crit ϕ_{max} was limited to: $\phi_{\text{max}} = \phi_{\text{ag}_{\text{max}}} \phi_{i_{\text{max}}} = 0.3$.

Fig. 2 illustrates spatial arrangements of RBCs for three internal hematocrits of 40%, 50%, and 60%. In these simulations, the aggregate radius had a constant value of $r_{\rm ag}/a = 6.32$ (i.e., $r_{\rm ag} = 17.39 \ \mu$ m) and a constant systemic hematocrit of 20%.



Fig. 2. Distributions of aggregated red blood cells for a constant aggregated radius $r_{ag} = 6.32a = 17.32 \ \mu\text{m}$ and a constant systemic hematocrit $\phi = 20\%$ at three internal hematocrits: (a) $\phi_i = 40\%$, (b) $\phi_i = 50\%$, and (c) $\phi_i = 60\%$. The displayed areas are $300 \times 300 \ \mu\text{m}$ to enhance the red blood cell visualization.

2) Distribution of Effective Particles for the Computation of S_{ag} With the EMTSFM: Once the effective particle radius was specified, particles were randomly distributed using a random number generator with non-overlapping positions to give the desired area fraction ϕ_{ag} . Several distributions were performed with different effective particle radii varying from 3.16a (= 8.69μ m) to 7.95a (= 21.86μ m) corresponding to different studied aggregate sizes.

B. BSC Computation

For each distribution of RBCs (or effective particles, respectively), density matrices D (or D_{ag}) were computed by dividing the square simulation plane L^2 in N_p^2 pixels (here, $N_p = 512$) and by counting the number of RBCs (or the number of effective particles) falling into each pixel. These matrices represented samplings of microscopic density functions defined by D(r) for the BSC computation with the SFM or by $D_{ag}(r')$ for the BSC_{eq} computation with the proposed EMTSFM:

$$D(\mathbf{r}) = \sum_{i=1}^{N} \delta(\mathbf{r} - \mathbf{r}_{i}),$$

$$D_{\rm ag}(\mathbf{r}') = \sum_{i=1}^{N_{\rm ag}} \delta(\mathbf{r}' - \mathbf{r}_{i}'),$$
(10)

where N is the number of RBCs in blood, $N_{\rm ag}$ is the number of aggregates, and δ is the Dirac distribution. According to (3) and (8), and as explicitly described in [14, Appendix], the structure factors can thus be described by

$$\begin{split} S(-2\mathbf{k}) &= E\left[\frac{1}{N} \left| \int D(\mathbf{r}) e^{-i2\mathbf{k}\mathbf{r}} \mathrm{d}\mathbf{r} \right|^2 \right] \text{ and } \\ S_{\mathrm{ag}}(-2\mathbf{k}) &= E\left[\frac{1}{N_{\mathrm{ag}}} \left| \int D_{\mathrm{ag}}(\mathbf{r}') e^{-i2\mathbf{k}\mathbf{r}'} \mathrm{d}\mathbf{r}' \right|^2 \right]. \end{split}$$
(11)

Each structure factor was thus computed by averaging 2-D fast Fourier transforms (2-D FFT) of 400 density matri-

ces for averaging purposes. The FFTs gave the structure factor values $S(-2\mathbf{k})$ and $S_{\rm ag}(-2\mathbf{k})$ on a centered grid of wavevectors between $\pm \pi N_p/2L$ with a step of $\Delta k = \pi/L$. For the SFM, the BSC_{SFM} was thus obtained using (1). For the EMTSFM, the BSC_{eq} was computed using (7).

IV. Results

In the following, the relative errors for BSC_{eq} computed with the new EMTSFM were evaluated as $(BSC_{eq} - BSC_{SFM})/BSC_{eq}$, where the BSC_{SFM} of the SFM is assumed to be the exact solution.

A. Influence of the Internal Hematocrit

A key feature of the new simulation method was the possibility to simulate randomly various internal hematocrits corresponding to different degrees of compactness of aggregates. Frequency-dependent backscattering coefficients with different internal hematocrits from 30% to 60% are given in Figs. 3(a) and 3(b) at systemic hematocrits of 10% and 20% over a large frequency range (4 to 100 MHz). Corresponding relative errors for the BSC_{eq} are given in Figs. 3(c) and 3(d). It was assumed that the aggregates had a constant radius of $r_{\rm ag}/a = 6.32$ (i.e., $r_{\rm ag} =$ 17.39 μ m). Note that for the 20% systemic hematocrit, internal hematocrits smaller than 40% could not be computed because the corresponding area fractions of aggregates were too high: $\phi_{ag} > 0.5$. As seen in Figs. 3(a) and 3(b), as the internal hematocrit ϕ_i increases, the BSC_{SFM} amplitude increases at low frequencies (<23 MHz) and decreases at high frequencies (>28 MHz). The first peaks of the BSC_{SFM} are between 18.0 and 20.6 MHz for all simulated conditions. The relative error for BSC_{eq} was less than 30% for frequencies below 18 MHz [see Figs. 3(c)and 3(d)]. This 18 MHz frequency corresponds to a product $kr_{\rm ag} = 1.27$. More generally, at frequencies less than 23 MHz, backscattering coefficients obtained with the SFM and the EMTSFM have the same behaviors: the first peaks of BSC_{SFM} and BSC_{eq} occur at the same frequencies



Fig. 3. (a) and (b) Dependence of the backscattering coefficients (BSCs) on the internal hematocrit: (a) $r_{\rm ag}/a = 6.32$ and $\phi = 10\%$, and (b) $r_{\rm ag}/a = 6.32$ and $\phi = 20\%$. The symbols are the BSC_{SFM} computed with the structure factor model (SFM) and the dashed lines are the BSC_{eq} computed with the effective medium theory combined with the SFM (EMTSFM). (c) and (d) Corresponding relative errors for the BSC_{eq} versus frequency.

and the magnitude of both $\rm BSC_{SFM}$ and $\rm BSC_{eq}$ increase as the internal hematocrit increases.

A quantitative ultrasonic parameter that has often been used for tissue characterization is the spectral slope (SS). The SS is the linear slope of the BSC as a function of frequency on a log-log scale. The variation of SS with the internal hematocrit is given in Fig. 4 for systemic hematocrits of 10% and 20%. The solid lines represent the SSs for the BSC_{SFM} computed with the SFM and the dashed lines represent the SSs for the BSC_{eq} computed with the EMTSFM. For both SFM and EMTSFM, the SS behavior versus the internal hematocrit is the same: the SS decreases with increasing internal hematocrit, except for the configuration $\phi = 20\%$, $\phi_i = 40\%$. Note also that the SS of the EMTSFM slightly underestimated the SS of the SFM with a maximum difference of 0.23.

B. Influence of the Aggregate Size

Fig. 5 shows backscattering coefficients as a function of frequency for different aggregate sizes and for systemic hematocrits of 10%, 20%, and 30%. In these simulations, the internal hematocrit was the highest: $\phi_i = 60\%$. The symbols represent the BSC_{SFM} computation for the disaggregated case $(r_{\rm ag}/a = 1)$ and for aggregated cases with radii of $r_{\rm ag}/a = 3.16$, 5.0, and 7.07. The corresponding



Fig. 4. Spectral slope in the frequency bandwidth 5 to 15 MHz as a function of the internal hematocrit at different systemic hematocrits, the aggregate size being fixed ($r_{\rm ag}/a = 6.32$). The solid lines are the spectral slopes (SSs) for the BSC_{SFM} computed with the structure factor model (SFM) and the dashed lines are the SSs for the BSC_{eq} computed with the effective medium theory combined with the SFM (EMTSFM).

 BSC_{eq} computed with the EMTSFM [see (7)] are given in dashed lines. For frequencies less than 20 MHz, the amplitudes of both BSC_{SFM} and BSC_{eq} increase with the size of aggregates. Moreover, the peaks of both BSC_{SFM} and BSC_{eq} occur at lower frequencies as the aggregate radius increases.

The low-frequency approximation of the EMTSFM (i.e., using the Percus-Yevick packing factor) was also studied for the same configurations. The BSC_{eq} values computed by (9) are given in Fig. 6, along with plots of the BSC_{SFM} computed with the SFM. As expected, the error for BSC_{eqLF} is larger than the error for BSC_{eq} at high frequencies. The error between BSC_{eqLF} and BSC_{SFM} increases as the systemic hematocrit ϕ increases. Nevertheless, it is interesting to observe that at low frequencies less than 5 MHz, the amplitude of the BSC_{eqLF} matches the exact BSC_{SFM} very well, as expected.

For systemic hematocrits of 10%, 20%, and 30%, the relative errors for BSC_{eq} and BSC_{eqLF} were studied for a constant value of the internal hematocrit $\phi_i = 60\%$ and for eleven sizes of r_{ag}/a varying from 3.16 to 7.95 with a step of around 0.5. For each case, the frequency limit f_1 and the corresponding product $k_l r_{ag}$ for which the relative error was less than 30% were determined. Results are summarized in Fig. 7. The solid lines are for the EMTSFM and the dashed lines for the low-frequency approximation of the EMTSFM using the Percus-Yevick packing factor. Fig. 7 reveals that the EMTSFM gives accurate estimates of BSC_{SFM} until $kr_{ag} \leq 1.32$ on average (i.e., for all the tested values of r_{ag}/a), whereas the EMTSFM at low frequency is only accurate until approximately $kr_{ag} \leq 0.73$.

Fig. 8 shows the influence of the aggregate size on the SS for different systemic hematocrits. The solid lines represent the SSs for the BSC_{SFM} computed with the SFM and the dashed lines represent the SSs for the BSC_{eq} computed with the EMTSFM. The results of SS presented in



Fig. 5. Dependence of the backscattering coefficients (BSCs) for different aggregate sizes and a constant internal hematocrit $\phi_i = 60\%$ at systemic hematocrits of 10%, 20%, and 30%. The symbols are the BSC_{SFM} computed with the structure factor model (SFM) and the dashed lines are the BSC_{eq} computed with the effective medium theory combined with the SFM (EMTSFM).



Fig. 6. Dependence of the backscattering coefficients (BSCs) for different aggregate sizes and a constant internal hematocrit $\phi_i = 60\%$ at systemic hematocrits of 10%, 20%, and 30%. The symbols are the BSC_{SFM} computed with the structure factor model (SFM) and the dashed lines are the BSC_{eque} computed with the effective medium theory combined with the SFM (EMTSFM) using the Percus-Yevick approximation.

Fig. 8 showed that the backscattered power by disaggregated RBCs ($r_{\rm ag}/a = 1$) increased with the third power of wave frequency (SS = 3), as was expected using the Rayleigh theory in 2-D. The SS behavior with the aggregate size is the same for both SFM and EMTSFM: as the aggregate size increases, the SS decreases at $\phi =$ 10%, whereas the SS increases at $\phi = 20\%$ and 30%. The maximum difference between slopes of the EMTSFM and slopes of the SFM is around 0.35 for all studied hematocrits (i.e., relative error for the SS of BSC_{eq} around 9%) and it corresponds to the largest aggregate size.

V. DISCUSSION

A. Limitation of the EMTSFM With Respect to kr_{aa}

Based on the new EMTSFM theory, the BSC from aggregated RBCs was parameterized by three indices: the aggregate size, internal hematocrit, and systemic hematocrit. The aim of our study was to estimate the goodness of fit of the EMTSFM approximation in comparison with the SFM. We found that the EMTSFM had a relative error of less than 30% for a mean product $kr_{\rm ag} \leq 1.32$ at all internal hematocrits and systemic hematocrits studied. Although the EMTSFM gave estimates with large errors for $kr_{\rm ag} > 1.32$, the EMTSFM fitted very well the BSC_{SFM} curve computed with the SFM for all tested factors (i.e., $r_{\rm ag}, \phi_i$ and ϕ) in the frequency bandwidth between 4 MHz and the frequency position of the first peak. This result is qualitatively consistent with works in optics [24], in which the limits of applicability of the interference approximation based on the structure factor were found for a volume fraction of particles smaller than 10% and for a product kR < 3.5 (or kR < 1.5, respectively) when the relative refractive index of the particles is 1.19 (or 1.8). These limits were determined by comparing theoretical predictions of the interference approximation with theoretical predictions of a much more exact quasi-crystalline approximation and with experimental data of a polystyrene latex suspension in water. In comparison with our results in acoustics in which the relative acoustic impedance of



Fig. 7. (a) Frequency limit $f_{\rm l}$ for which the relative error of BSC_{eq} was less than 30% as a function of the aggregate size studied. The solid lines are the effective medium theory combined with the structure factor model (EMTSFM) and the dashed lines are the low-frequency approximation of the EMTSFM. (b) Corresponding product $k_{\rm l}r_{\rm ag}$.

the RBCs is around 1.12, the limit of applicability of the proposed EMTSFM in terms of particle size is surprisingly close to the optic limit $kR \leq 1.5$, when one considers a large relative refractive index of 1.8. In terms of particle concentration, the EMTSFM showed no limitation with respect to the studied hematocrit (up to 30%), contrary to the interference approximation, valid up to a volume fraction of 10%.

B. Variation of the Spectral Slope With ϕ_i and With r_{ag}

On the goodness of the EMTSFM modeling the SS, notice that the variations of SS with ϕ_i and r_{ag} followed similar trends for both SFM and EMTSFM (Figs. 4 and 8). The SS of the EMTSFM slightly underestimated the SS of the SFM with a maximum relative error of 9%. As can be observed in Fig. 8, as the aggregate size increased, the difference between slopes of the EMTSFM and SFM increased. This is due to the constant frequency bandwidth chosen between 5 and 15 MHz for all studied aggregate sizes. Indeed, in the frequency bandwidth between 4 MHz and the frequency position of the first peak, the relative error for



Fig. 8. Spectral slope in the frequency bandwidth 5 to 15 MHz as a function of the aggregate size at different systemic hematocrits, the internal hematocrit being fixed ($\phi_i = 60\%$ for aggregated red blood cells). The solid lines are the spectral slopes (SSs) for the BSC_{SFM} computed with the structure factor model (SFM) and the dashed lines are the SSs for the BSC_{eq} computed with the effective medium theory combined with the SFM (EMTSFM).

the BSC_{eq} increased with the frequency as was shown in Figs. 3(c) and 3(d). Because the relative error was less than 30% for a mean product $kr_{\rm ag} \leq 1.32$, the mean relative error in the frequency bandwidth 5 to 15 MHz for the smallest aggregates was smaller than the relative error for the largest aggregates in the same frequency bandwidth.

Contrary to blood modeling, the ultrasound characterization of tissues such as the breast [4] or the liver [25] assumes scatterers to be randomly distributed at a low number density. Tissue models are generally based on 3-D spatial autocorrelation functions describing the shape and isotropic random distribution of scatterers in the medium, and modeling of the spatial autocorrelation function could be Gaussian, spherical, or exponential [22], [25]. Based on these models, the SS is an indicator of the scatterer size, and a decrease in slope usually corresponds to an increase in scatterer size [25]. That is why it is interesting to discuss the variation of the SS with the aggregate size in the framework of the SFM or EMTSFM which consider the structure factor (see Fig. 8). For the lowest systemic hematocrit of 10%, the SS decreased with the increase in aggregate size. Because the hematocrit is low, the known behavior of the SS was observed. However, the SS increased with the aggregate size for systemic hematocrits of 20 and 30%. The structure factor, which models the spectral behavior of the scatterer spatial distributions, caused this behavior of the SS. Indeed, according to (1) [or (7), respectively], the BSC_{SFM} frequency dependence (or the BSC_{eq} frequency dependence) is determined by the frequency dependences of σ_b and S (or σ_{ag} and S_{ag}). To illustrate this, the structure factors S and S_{ag} corresponding, respectively, to the SFM and the EMTSFM are displayed in Fig. 9 for a fixed systemic hematocrit $\phi = 30\%$, a fixed internal hematocrit $\phi_i = 60\%$, and three aggregate radii $r_{\rm ag}/a = 3.16, 5.0, \text{ and } 7.07.$ The plot of S is represented in a larger frequency bandwidth up to 320 MHz to show the high-frequency oscillations of the structure factor around 1. In the frequency bandwidth 5 to 15 MHz, the frequency dependence of the structure factors S and S_{ag} do not follow a simple power law, because the structure factors Sand $S_{\rm ag}$ versus frequency on a log-log scale are non-linear curves. One could observe that, in the frequency bandwidth 5 to 15 MHz, both the structure factors S and S_{ag} increase with the frequency, and the larger the aggregate size is, the larger is the increase. That is why the SS for both BSC_{SFM} and BSC_{eq} increased above 3 with the aggregate size for this studied systemic hematocrit of 30%. The increase of SS with the aggregate size we observed in the current study is consistent with an earlier 2-D numerical simulation study performed by Fontaine and Cloutier [17]. In this 2-D study, the SS between 5 and 25 MHz could increase from 3.1 to 3.3 when the level of aggregation increased for a hematocrit of 40%. Moreover, in vitro Couette flow experiments performed in [18] showed the SS greater than 4 between 9 and 15 MHz for shear rates of 2 and 10 s^{-1} at a systemic hematocrit of 40% [18, Fig. 4(b)].

It is important to note that we found that the SS decreased as ϕ_i increased for all studied systemic hematocrits. On the other hand, the behavior of the SS as a function of the aggregate size depended on the systemic hematocrit. As a consequence, the parameter SS seems well adapted to observe a change in the internal hematocrit. However, in experimental conditions where effects of aggregate size and internal hematocrit are mixed, the SS cannot be a useful index for blood characterization.

C. Computation of RBC Distributions

The method we proposed here to obtain the RBC spatial distribution did not take into consideration realistic interactions between RBCs. It was a simple and fast method to generate samples containing non-overlapping, identical, and isotropic aggregates. The main advantage of this method was the possibility to have various internal hematocrits with the same size of aggregates. The simulation results showed that the frequency position of the BSC first peak was very little affected by changes in the internal hematocrit, whereas it was greatly affected by changes in the aggregate size. Indeed, the change in frequency position was only 1.5 MHz (and, respectively, 2.1 MHz) for the hematocrit of 10% (and 20%) when the internal hematocrit increased from 30to 60%. Moreover, theoretical predictions of the BSC with the SFM showed that its amplitude increases as $r_{\rm ag}$ and ϕ_i increase (see Figs. 3 and 5). To our knowledge, the influence of the internal hematocrit on the BSC has not been studied previously, mainly because of the simulation methods used to realize distributions of aggregating RBCs. These methods were based on particle dynamics or statistical mechanics [12], [13], [15]–[17] and had the objective of mimicking the rheological behavior of blood. The RBC distributions obtained showed aggregates with anisotropic shape [12], [13], [15], [17], [26] and a polydispersity in terms of aggregate sizes, aggregate shapes, and internal hematocrits [12], [13], [15]–[17]. As mentioned previously, contrary to those earlier studies, the 2-D simulation method we proposed for distributing RBCs allowed isolation of the effects of the aggregate size and internal hematocrit to better understand the role of each parameter. In that sense, our study has provided some insights into the influence of the aggregate size and internal hematocrit on the BSC frequency dependency. The internal hematocrit and the aggregate size can greatly influence the BSC amplitude. The frequency position of the BSC first peak was found not to be significantly affected by changes in the internal hematocrit, whereas it was greatly affected by changes in the aggregate size.

D. On the Use of the EMTSFM In Vivo

Although the present work allowed a fundamental study on the influences of the internal hematocrit and aggregate



Fig. 9. (a) Structure factors S versus frequency used in the computation of the structure factor model (SFM) for a fixed systemic hematocrit $\phi = 30\%$, a fixed internal hematocrit $\phi_i = 60\%$, and three aggregate radii $r_{ag}/a = 3.16$, 5.0, and 7.07. (b) Structure factors S_{ag} versus frequency used in the computation of the effective medium theory combined with the SFM (EMTSFM) for the same aggregating conditions.

size on the BSC, the method we chose to distribute particles was limited to area fractions of aggregates $\phi_{\rm ag} \leq 0.5$, area fraction of RBCs within aggregates $\phi_i \leq 0.6$, and systemic hematocrits $\phi \leq 0.3$. One might thus question the use of the EMTSFM for *in vivo* experiments with a physiological hematocrit ϕ typically varying between 0.3 and 0.5. It is important to recall that the only intensive computation with the proposed EMTSFM is the structure factor S_{ag} , which needs the computation of effective particle distributions. If one considers the application of the EMTSFM in vivo, 3-D effective particle distributions must be considered as well as the range of typical volume fractions. For example, the volume fraction of RBCs within aggregates ϕ_i may physiologically be larger than 60% because RBCs are deformable. By considering a systemic hematocrit of 40% (and 20%, respectively) and extrema values of ϕ_i between 60 and 90%, the volume fraction of aggregates ϕ_{ag} would then vary between 44 and 66% (between 22 and 33%). For future simulations, particle distributions in 3-D could be easily generated using a random number generator up to a volume fraction of approximatively 0.3, meaning that the method used to compute the BSC_{eq} in this paper may easily be applied in 3-D up to a volume fraction ϕ of 20%. For such low volume fractions of cells, one could envisage the use the EMTSFM for other biomedical applications such as cancer [4], [27], where the cells can be locally densely packed (see, for example, [4, Fig. 5]). In the case of blood applications, to deal with a physiological hematocrit of 40% and a maximum volume fraction of aggregates of 66%, one may envisage distributing particles in a random close packing configuration in which the attainable volume fraction could be up to 64% [28].

On the use of the EMTSFM in vivo, one can also question the practical value of that model assuming isotropic aggregates. In human blood, low shear rates can promote the formation of RBC aggregates having anisotropic (i.e., rouleaux) or isotropic (i.e., clump) structures. The rouleaux-like pattern is typically associated with normal blood. However, as the binding energy between RBCs increases with inflammation [29], aggregates form clump structures such as in diabetes mellitus [30], [31]. The assumption of isotropic aggregates in the EMTSFM is thus valid as far as we are concerned with the study of pathological states. In the case of normal human rouleaux of RBCs, if the EMTSFM is applied to estimate structural parameters such as the internal hematocrit and the aggregate size, this assumption would obviously create a bias against the parameter estimation.

Another important aspect to consider is the assumption of identical aggregates in the current EMTSFM implementation. Under *in vivo* conditions with ultrasound measurements on a blood vessel, the shear rate distribution varies with the radial position, and consequently, the aggregate size distribution does too. That is why the back-scattered echoes from blood are generally analyzed over a rectangular or a Hamming window which is moved along the RF signal to examine the whole vessel at different depths [19], [32]. For example, at a central frequency of 25 MHz, the window length was typically around 400 μ m

[19], [32], [33]. RBC aggregates may thus be assumed to be locally identical. However, one may also expect that the aggregate size varies around a mean value for a given radial position within a blood vessel. An interesting simulation study was recently performed in 2-D by Vlad *et al.* [34]. According to this paper, the increase in particle size variance (or equivalently, in aggregate size variance) is expected to increase the BSC amplitude and not to significantly affect the frequency position of the BSC first peak [34, Fig. 2]. One could thus predict the bias against the parameter estimation with the EMTSFM if there is a variance in aggregate sizes. The discussion next considers small and large aggregates with respect to the observation of Vlad *et al.* [34].

In the case of small aggregates, the BSC first peak occurs at high frequencies and is thus generally not captured in the measured frequency bandwidth of the transducer (typically from 12 to 38 MHz for a 100% bandwidth at 25 MHz). Readers may refer to [18, Fig. 4(b)] and to [19, Fig. 3] to see measured BSCs from aggregating porcine RBCs in a controlled Couette flow. Because of the predicted increase in BSC amplitude with the increase in particle size variance [34], the assumption of identical aggregates in the current study may result in overestimations of the aggregate size and of the internal hematocrit. On the other hand, in the case of large aggregates, the BSC first peak occurs at low frequencies and would likely be within the measured frequency bandwidth of the transducer. In this case, the assumption of identical aggregates may not greatly affect the aggregate size estimation but may result in an overestimation of the internal hematocrit. The EMTSFM is thus expected to better estimate the size of large aggregates and the estimation of the internal hematocrit may be overestimated in the case of polydisperse aggregates, whatever their dimensions.

VI. CONCLUSION

The SFM is recognized to be a good model to simulate the BSC from aggregated RBCs. However, it cannot directly be applied to estimate the structural aggregate parameters in the framework of an inverse problem formulation. The EMTSFM was thus proposed to approximate the SFM and the goodness of fit of the EMTSFM in comparison with the SFM was examined. Contrary to the SFM, the EMTSFM can be implemented to estimate structural aggregate parameters.

The EMTSFM parameterizes the BSC by three indices: the aggregate size, the internal hematocrit (or aggregate compactness), and the systemic hematocrit. In comparison with the SFM, the EMTSFM provided accurate quantitative estimates of the BSC for $kr_{\rm ag} \leq 1.32$ for the studied conditions. Moreover, even if $kr_{\rm ag} > 1.32$, the EMTSFM provided a good approximation of the BSC: indeed, the BSC amplitude was well approximated in the frequency bandwidth between 4 MHz and the frequency position of the first peak, and the high-frequency positions of peaks were well identified. Another important finding was the variation of the SS with ϕ_i and r_{ag} . The SSs for both BSC_{SFM} and BSC_{eq} computed with the SFM and EMTSFM followed the same trends of variations with ϕ_i and r_{ag} . We found that the SS decreased as the internal hematocrit increased for all studied systemic hematocrits. On the other hand, the behavior of the SS as a function of the aggregate size depended on the systemic hematocrit. For the largest hematocrits (20 and 30%), the SS increased as the aggregate size increased. Consequently, the SS cannot be a useful index for blood characterization of RBC aggregation.

To conclude, the EMTSFM approximates the SFM satisfactorily for a product $kr_{\rm ag} \leq 1.32$. This suggests that the EMTSFM is an adequate model for blood characterization. An important contribution of this new model is the parameterization of the BSC with the aggregate compactness, which is a structural parameter not available in any other modeling strategies proposed in quantitative ultrasound imaging. In the future, it would be interesting to study two other important factors: the polydispersity in terms of aggregate sizes and internal hematocrits, and the non-sphericity of the aggregates. Another important study would be the estimation of structural aggregate parameters with the EMTSFM that could be compared with existing methods such as the SFSE [16], [18], [19] and the traditional Gaussian model developed by Lizzi [6], [25].

Appendix

The form factor is generally given in 3-D [21], [22]. That is why the demonstration to obtain the 2-D form factor of a fluid infinite cylinder of radius a is given here. In the Born approximation (i.e., for weak scattering conditions), the form factor has a simple expression that is related to the spatial Fourier spectrum of the impedance contrast (see [21, Eq. (5)] for the form factor expression in 3-D):

$$F(-2k,a) = \frac{1}{A_p^2 \gamma_z^2} \left| \int_{A_p} \gamma_z(\mathbf{r}) e^{2i\mathbf{k}\mathbf{r}} \mathrm{d}^2 \mathbf{r} \right|^2.$$
(12)

We can introduce cylindrical coordinates to obtain

$$F(-2k,a) = \frac{1}{A_p^2 \gamma_z^2} \left| \int_0^a \int_{-\pi}^{\pi} \gamma_z e^{2ikr \cos(\theta)} r \,\mathrm{d}r \,\mathrm{d}\theta \right|^2$$

$$= \frac{1}{A_p^2} \left| \int_0^a 2\pi r J_0(2kr) \,\mathrm{d}r \right|^2$$

$$= \frac{(2\pi)^2}{(\pi a^2)^2} \left(\frac{2ka J_1(2ka)}{(2k)^2} \right)^2$$

$$= \left(\frac{J_1(2ka)}{ka} \right)^2,$$
 (13)

where J_0 is the Bessel function of the first kind of order 0.

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Emilie Franceschini received her M.S. degree in mechanical engineering from the Ecole Supérieure d'Ingénieurs de Marseille in 2003, and her Ph.D. degree in acoustics from the University of Provence, Marseille, 2006. In 2007, she was a Postdoctoral Fellow at the Laboratory of Biorheology and Medical Ultrasonics, Research Center of the University of Montreal Hospital, Montreal, QC. Since October 2008, she has been a Researcher at the French National Centre for Scientific Research (CNRS) in the Laboratory of Mechanics

and Acoustics CNRS–UPR 7051, Marseille. Her current research interests include ultrasound imaging for biomedical applications, inverse problems, and ultrasound characterization of biological tissues at the microscopic level. She is a member of the French Acoustical Society (SFA) and IEEE.



Bloen Metzger received his M. S. degree from the Ecole Normale Supérieure de Lyon in 2003, and his Ph.D. degree in physics from the University of Provence, Marseille in 2006. In 2007, he was a Postdoctoral Fellow at the Benjamin Levich Institute, New York, NY. Since October 2007, he has been a Researcher at the French National Centre for Scientific Research (CNRS) in the laboratory IUSTI CNRS–UMR 6595, Marseille. His current research interests include microhydrodynamics, suspension flows at low Reynolds num-

bers, hydrodynamic diffusion, and heat and mass transfers in sheared suspensions.



Guy Cloutier (S'89–M'90–SM'07) obtained his B.Eng. degree in electrical engineering in 1984, and his M.Sc. and Ph.D. degrees in biomedical engineering in 1986 and 1990, respectively. Between 1990 and 1992, he was a post-doctoral fellow at The Pennsylvania State University with Prof. K. Kirk Shung. Prof. Cloutier is Director of the Laboratory of Biorheology and Medical Ultrasonics at the University of Montreal Hospital Research Center (http://www.lbum-crchum.com), and Professor of Radiology and Biomedical Engi-

neering at the University of Montreal. His research interests are in quantitative ultrasound imaging of red blood cell aggregation; quasi-static and dynamic ultrasound elastography of atherosclerotic plaques, vascular aneurysms, deep vein thrombi, and breast cancers; 3-D morphologic and hemodynamic assessment of lower limb arterial stenoses; and mathematical and biomechanical modeling. He has published more than 120 peer-reviewed articles in these fields, holds 8 patents, and was recipient of the National Scientist award of the Fonds de Recherche en Santé du Québec (2004–2009).