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Abstract. Diffuse correlation spectroscopy (DCS) measurements of blood flow rely on the sensitivity of the temporal autocorrelation function of diffusively scattered light to red blood cell (RBC) mean square displacement (MSD). For RBCs flowing with convective velocity \( v_{RBC} \), the autocorrelation is expected to decay exponentially with \( e^{-\mu t} \), where \( t \) is the delay time. RBCs also experience shear-induced diffusion with a diffusion coefficient \( D_{\text{shear}} \), and an MSD of \( 6D_{\text{shear}} \). Surprisingly, experimental data primarily reflect diffusive behavior. To provide quantitative estimates of the relative contributions of convective and diffusive movements, we performed Monte Carlo simulations of light scattering through tissue of varying vessel densities. We assumed laminar vessel flow profiles and accounted for shear-induced diffusion effects. In agreement with experimental data, we found that diffusive motion dominates the correlation decay for typical DCS measurement parameters. Furthermore, our model offers a quantitative relationship between the RBC diffusion coefficient and absolute tissue blood flow. We thus offer, for the first time, theoretical support for the empirically accepted ability of the DCS blood flow index \( (BF_i) \) to quantify tissue perfusion. We find \( BF_i \) to be linearly proportional to blood flow, but with a proportionality modulated by the hemoglobin concentration and the average blood vessel diameter.

Keywords: diffuse correlation spectroscopy; blood flow; Monte Carlo; photon migration.

1 Introduction

Diffuse correlation spectroscopy (DCS) is increasingly being used as a noninvasive method to quantify tissue blood flow, especially for neurological applications. DCS measurements quantify the temporal autocorrelation of fluctuations in the intensity of diffusely reflected light that has traveled through tissue. These fluctuations are almost entirely driven by the motion of red blood cells (RBCs) and thus by blood flow. Although DCS blood flow measurements have been validated against several alternative perfusion quantification methods, a discrepancy remains between the expected nature of RBC displacement and experimental results. Specifically, instead of an expected ballistic random walk with a uniform spatial velocity distribution, RBC motions measured by DCS have the characteristics of diffusive motion.

Previous publications from our group as well as Ninck et al. hypothesized that DCS measurements are also sensitive to the shear-induced diffusion of RBCs in blood flow, a phenomenon previously observed through particle tracking experiments. Shear-induced diffusion refers to the random walk-like motion of particles in a shear flow due to interparticle hydrodynamic interactions. Although it was noted that the effective RBC diffusion coefficient is proportional to the flow shear rate, the question of what are the relative contributions of diffusive and convective motions to the observed intensity autocorrelation decay has not been addressed previously.

In this publication, we seek to answer this question using Monte Carlo simulations of light scattering through tissues with varying densities and diameters of vessels. We explicitly model the location of each light scattering event within the vessel flow profiles and consider both convective motion and the corresponding shear-induced diffusion. Through these simulations, we obtain quantitative estimates of the contributions of both processes to the decay of the intensity temporal autocorrelation function. We present definitive evidence that diffusive motion dominates the decay for typical DCS experimental conditions and derive explicit expressions for the relationship between the measured RBC diffusion coefficient and absolute blood flow. Furthermore, we show that the commonly used DCS blood flow index \( BF_i \) derived using the correlation diffusion equation (CDE) for a semi-infinite geometry is directly proportional to absolute blood flow, but this proportionality is modulated by the hematocrit and the average vessel diameter in the volume of tissue probed by light.

2 Theory and Methods

2.1 Correlation Diffusion Equation

DCS measurements of blood flow are generally analyzed using the CDE. The CDE is derived from the correlation transfer equation (CTE) under the assumption that the probability of light scattering is much greater than the probability of light absorption, i.e., reduced scattering coefficient \( \mu_i \) is much greater than the absorption coefficient \( \mu_a \). The CTE permits calculation of the electric field temporal autocorrelation...
function $G_1(\tau)$ under more general conditions of photon migration. The CTE is similar to the traditional radiative transfer equation (RTE), which describes the propagation of light intensity through scattering media. The difference between the CTE and the RTE is that the CTE describes the propagation of the time-varying specific intensity, which represents an angular spectrum of the mutual coherence function. The temporal autocorrelation function is obtained as an integral of the time-varying specific intensity over all solid angles.

Generally, an analytic solution of the CDE for a semi-infinite medium is used to fit the experimental measurements, i.e.,

$$G_1(\rho, \tau) = \frac{3\mu'_s}{4\pi} \left[ \exp(\sqrt{K^2(\tau)}) - \exp(K_0^2(\tau)) \right],$$

where

$K^2 = 3\mu'_s + \mu_s^2 k_0^2 \exp(\Delta r^2(\tau)),$

where $\mu'_s$ is the reduced scattering coefficient, $\mu_s$ is the absorption coefficient, $k_0$ is the wavenumber of light in the scattering medium, $\alpha$ is the probability of scattering from a moving scatterer, $\rho$ is the source–detector separation, $r_1 = (\rho^2 + z_1^2)^{1/2}$, $r_2 = (\rho^2 + z_2^2)^{1/2}$, $z_0 = \mu_s^{-1}$, and $z_h = 1.76/\mu'_s$, given an index of refraction for tissue of 1.37.

It is customarily expected that the scatterer motion measured by DCS arises almost exclusively from RBCs. This is justified by experimental studies in animals, which showed that cessation of blood flow results in a nearly 100-fold reduction in the observed dynamics. In this paper, we will assume that RBCs of blood flow results in a nearly 100-fold reduction in the dynamic scattering events, and we will use $P_{RBC}$ instead of $\alpha$ to denote the probability of scattering from an RBC.

The mean square displacement (MSD) $\langle \Delta r^2(\tau) \rangle$ of the RBCs is generally given by

$$\langle \Delta r^2(\tau) \rangle = 6D\tau + v_{RBC}^2 \tau^2,$$

where $\tau$ is the delay time, $D$ is the diffusion coefficient, and $v_{RBC}$ is the speed of the RBCs. Note that Eq. (3) can be expanded to use the Langevin formulation for diffusion, 13,19,20 accounting for the diffusive motion velocity vector randomization that occurs at short-time delays.

### 2.2 Monte Carlo Simulations

Investigations with DCS have generally assumed that the motion of tissue scatterers is uncorrelated. Specifically, the assumption is that the phase shifts accumulated by the motions of scattering particles is uncorrelated, which depends on the motion of the scattering particle and the scattering angle. This assumption is generally valid when photons are scattering from particles undergoing Brownian motion. In the case of moving RBCs, this assumption is valid if the photon direction is randomized between scattering events from RBCs. This is true if the scattering is isotropic (i.e., the average cosine of the scattering angle is $g = 0$) or if the photons scatter no more than once inside a vessel and the direction is randomized before scattering from another RBC in another vessel. The optical properties of blood at 800 nm indicate, 21 however, that for a typical hematocrit of 40%, $g \approx 0.98$ and the scattering length $l_s^2 \approx 12$ $\mu$m, whereas the absorption length $l_a$ is over 250 times larger, at around 3.3 mm. Thus, photons entering any blood vessel larger than a capillary will most likely undergo multiple consecutive scattering events within the vessel with correlated linear motions of the RBCs. This breakdown of the uncorrelated motion assumption raises the possibility that the typically used solutions of the CTE, and thus the CDE, are not good models for analyzing DCS measurements of blood flow.

To properly account for the effect of correlated motions on the decay of $G_1(\tau)$, we use Monte Carlo simulations of photon migration through highly scattering dynamic media and record details of the scattering angles for consecutive scattering events within vessels. Our Monte Carlo code is derived from what Boas originally utilized in Ref. 16 based on earlier work by Middleton and Fisher, 22 Durian, 23 and Koelink et al. 24 The original Monte Carlo simulation propagates photons according to the scattering length and scattering anisotropy of the tissue, which we describe in more detail in Ref. 25. Photons are launched from a source position and recorded at discrete detector locations. For each detected photon, the total path length $L$ is recorded as well as the total dimensionless momentum transfer accumulated over all scattering events $Y$. The momentum transfer for a photon scattering event is given by $q = \hat{k}_\text{out} - \hat{k}_\text{in}$, where $\hat{k}_\text{out}$ and $\hat{k}_\text{in}$ are the photon wavevectors scattered from and incident on the scattering center, respectively, and the total dimensionless momentum transfer is given by $Y = \sum q^2/2k_0^2$, where the sum is over scattering events. The temporal field autocorrelation function is then calculated by

$$G_1(\tau) = \frac{1}{N_p} \sum_{n=1}^{N_p} \exp \left( -\frac{1}{3} Y_n k_0^2 \langle \Delta r^2(\tau) \rangle \right) \exp(-\mu_s L_n),$$

where $N_p$ is the number of photons detected, $Y_n$ is the total dimensionless momentum transfer for photon $n$, $L_n$ is the total path length of photon $n$, $\mu_s$ is the absorption coefficient, and $\langle \Delta r^2(\tau) \rangle$ is the MSD of the scattering particles at a time delay of $\tau$.

The principal advance in the Monte Carlo simulations used in this paper is the introduction of blood vessels with different optical and dynamic properties than the surrounding tissue and with specified internal flow profiles, as shown in Figs. 1 and 2. Photon migration is simulated in a semi-infinite medium with the air tissue interface in the $(x, y)$ plane at $z = 0$. The blood vessels share a common radius $\alpha$ and are all oriented along the $y$-axis with an even spacing in $x$ and $z$ of $h_{space}$. The resulting translational symmetry allows for significant savings in the amount of memory needed to represent the tissue structure for the Monte Carlo simulation. We have verified that this preferred orientation of the vessels only introduces a small bias in our results compared to a truly random orientation of vessels, as expected, given that we typically simulate photons that have propagated a net-distance of several centimeters and have scattered hundreds of times.

Photons are launched at a point on the surface along the $z$-axis and are detected through a 2-mm-diameter circular aperture, a distance $\rho$ from the source point. Figure 2 shows a segment of a photon path and the parameters we record during the photon propagation. For each scattering step during the simulation, we explicitly test for the photon crossing the boundary of a cylindrical vessel. If a crossing occurs, the remaining propagation length for the scattering step is renormalized based on the new scattering coefficient as we describe in Ref. 25. For each detected photon, we record for every scattering event inside a cylindrical vessel the $\rho$, $\Delta \rho$, and $\phi$. The average length $L_\text{avg}$ is calculated as $L_\text{avg} = \sum L_n$, where $L_n$ is the total path length of photon $n$.
vessel the index of the vessel, the radial position of the scattering event, the magnitude of the momentum transfer $|\mathbf{q}|$, and the component of $\mathbf{q}$ along the vessel direction ($y$-axis). Recording the vessel index enables us to determine consecutive scattering events within a vessel. Knowing the radial position of the scattering event permits us to consider a radial distribution for the convective and diffusive dynamics of the RBCs. We record the $y$ component of $\mathbf{q}$ as it is needed to consider the correlated convective motion of the RBCs along the $y$-axis. All Monte Carlo simulations in this paper use a source–detector separation of 2 cm, an $(x,z)$ plane cross section of $6 \times 3$ cm, and make use of translational symmetry to represent an infinite extent in $y$.

Each simulation run launches $10^8$ photons.

To calculate the temporal field autocorrelation function, we start with the definition of $G_1(\tau)$ and expand it to sum over all detected photons

$$G_1(\tau) = \langle E(0) E^*(\tau) \rangle$$

$$= \frac{1}{N_p} \sum_{n=1}^{N_p} \left\langle \exp \left( i \sum_{m=1}^{N_{\text{sc}}(n)} \mathbf{q}_{n,m} \cdot \mathbf{\Delta r}_{n,m}(\tau) \right) \right\rangle \times \exp \left( -\sum_{p=1}^{N_{\text{in}}} \mu_{n,p} L_{n,p} \right). \quad (5)$$

Note that one generally reports the normalized temporal field autocorrelation function $g_1(\tau)$, which is just $G_1(\tau)$ normalized by the average intensity. The angle brackets $\langle \rangle$ indicate the ensemble average over scattering angles and particle displacements. The number of photons received at the specific detector is $N_p$. The number of scattering events for the $n$'th detected photon is $N_{\text{sc}}(n)$, where $\mathbf{q}_{n,m}$ is the momentum transfer for the $m$'th scattering event of the $n$'th photon and $\mathbf{\Delta r}_{n,m}(\tau)$ is the
displacement of the corresponding scattering particle at time lag \( \tau \). The total path length of the \( n \)th photon in the \( p \)th tissue type is given by \( L_{n,p} \) and \( \mu_{a,p} \) is the absorption coefficient of the \( p \)th tissue type.

We are considering the convective and shear-induced diffusive movement of RBCs such that the MSD of the RBCs is given by Eq. (3). Although the strength of the diffusive motion correlates with the magnitude of the local shear, the actual diffusive displacements are spatially uncorrelated with the convective displacements. Therefore, note that \( q_{n,m} = \Delta r_{n,m}(\tau) \) is generally much smaller than 1, and Eq. (5) can be rewritten as

\[
G_1(\tau) = \sum_{n=1}^{N} \left[ \frac{1}{2} \sum_{m=1}^{N} \left( q_{n,m} + \frac{\tau}{\lambda_{FS}} \right) \cdot v_{n,m} \right]^2 \exp \left( \sum_{m=1}^{N} \mu_{a,p} \tau_{n,m} \right) \exp \left( \sum_{p=1}^{N} \mu_{a,p} \tau_{n,p} \right).
\]

This formulation reveals the expected decay of \( G_1(\tau) \) due to the RBC displacements arising from diffusion and convective flow. Note that the decay arising from flow speed accounts for the correlated RBC displacements as the sum \( (\sum_{n=1}^{N} \sum_{m=1}^{N} q_{n,m} + \frac{\tau}{\lambda_{FS}} \cdot v_{n,m})^2 \) is performed over the \( N_{n,m} \) consecutive scattering events of the \( n \)th photon in the \( i \)th vessel. The sum over the \( N_{n,m} \) vessels entered by the \( n \)th photon is outside the \( \dots \) because the phase shifts \( q_{n,m} \cdot \frac{\tau}{\lambda_{FS}} \cdot v_{n,m} \) are assumed uncorrelated between vessels, as the scattering angle and flow speed vector are generally randomized between photon interactions with different vessels. All of the information for calculating Eq. (6) is recorded by our Monte Carlo simulation. Given details about the RBC speed and shear-induced diffusion, we can now explicitly test which dominates the decay of \( G_1(\tau) \) and whether the CDE provides a good model fit and allows estimation of blood flow.

### 2.3 Red Blood Cell Laminar Flow Profile and Shear-Induced Diffusion

Our Monte Carlo simulations permit us to incorporate radial distributions for the flow speed and diffusion coefficient. A general form of the flow speed profile in vessels has been proposed to be

\[
v_{RBC}(r) = v_{\text{max}} \left[ 1 - \left( \frac{r}{R} \right)^a \right],
\]

where \( v_{\text{max}} \) is the maximum speed at the center of the vessel, \( R \) is the radius of the vessel, \( a \) is a scale factor that allows for a nonzero velocity at the wall, and \( k \) determines the degree of ”bluntness” of the profile. In general, the smaller the vessel the more blunt the profile. For sufficiently large vessels, the flow speed profile can be considered to be laminar; thus, in our simulations, we only consider parabolic flow profiles with \( a = 1 \) and \( k = 2 \). We believe this assumption captures the essence of the influence of the flow profile on RBC dynamic scattering events. If needed, other \( a \) and \( k \) values can be used in the future to account for the various degrees of flow profile blunting that have been observed in blood microcirculation investigations.

RBC dynamics, while flowing within a vessel, have been studied in numerous rheological studies including video microscopy to track the motion of RBCs ex vivo\(^{15} \) and whole blood in microchannels,\(^{17} \) as well as in vivo in rat venules.\(^{18} \) These studies have revealed that shear induces a diffusive behavior in the movement of the RBCs with the diffusion coefficient proportional to the shear rate, i.e.,

\[
D(r) = \alpha_{\text{shear}} v(r) = \alpha_{\text{shear}} \frac{\partial v_{RBC}}{\partial r} = 2\alpha_{\text{shear}} \frac{v_{\text{max}}^2 r}{R^2}. \tag{8}
\]

For our simulations, we use the value of \( \alpha_{\text{shear}} = 10^{-6} \text{ mm}^2/\text{ s} \) from Goldsmith and Marlow.\(^{15} \)

Of note, a prior study by Wu et al.\(^{29} \) has shown that the gradient in the speed of scattering particles flowing within a channel can be the leading factor in the decay of the autocorrelation function. This is an informative study, but the parameters used are quite different from those encountered in biological tissue, leading to negligible levels of shear-induced diffusion. This is due to the use of a single flow channel with a much smaller shear rate than that experienced by RBCs in blood vessels, as well as a much smaller particle concentration and scattering coefficient. Furthermore, this study did not consider multiple flow channels in an otherwise static scattering medium.

### 2.4 Relating the Diffuse Correlation Spectroscopy Blood Flow Index to Absolute Blood Flow

We are now in a position to perform Monte Carlo simulations to calculate \( g_{1,MC}(\tau) \) for RBCs flowing at varying speeds in vessels of varying diameters and spacings. We can then fit the result with \( g_{1,\text{CDE}}(\tau) \) obtained from the CDE to determine if the CDE remains a good model and to establish a quantitative relationship between the CDE blood flow index \( BF_{\text{CDE}} \) and the absolute blood flow \( BF_{\text{abs}} \). For a semi-infinite medium, we use \( g_{1,\text{CDE}}(\tau) \) given in Eq. (1) and estimate \( K \) from Eq. (2). The optical properties and wavelength of light are generally known in Eq. (2), leaving us to estimate \( BF_{\text{CDE}}(\Delta r^2(\tau)) \). We define the blood flow indices

\[
BF_{\text{conv}} = \sqrt{P_{\text{RBC}}(r^2)} \quad \text{and} \quad BF_{\text{diff}} = P_{\text{RBC}}(D)
\]

by fitting either the convective or diffusive MSD model to \( g_{1,\text{MC}}(\tau) \), where either \( BF_{\text{conv}} \) is expected to be linearly proportional to flow speed. The angle brackets for \( \langle v \rangle \) and \( \langle D \rangle \) are an explicit recognition that the \( BF_{\text{conv}} \) measures the average speed \( v \) or diffusion coefficient \( D \). Note that the probability of scattering from an RBC is not generally known a priori. In the context of photon diffusion theory, we would expect that \( P_{\text{RBC}} \) is given by the product of the volume fraction of blood, \( f_{\text{blood}} = \pi R^2 \mu_{\text{space}}^2 \), and the ratio of the blood to average tissue reduced scattering coefficient, \( \mu_{\text{space}}^\prime / \mu_{\text{avg}}^\prime \), i.e.,

\[
P_{\text{RBC}} = \frac{\pi R^2}{\mu_{\text{space}}^\prime / \mu_{\text{avg}}^\prime}.
\]

The average tissue-reduced scattering coefficient is given by the volume fraction weighted average of the tissue and blood coefficients \( \mu_{\text{avg}} = f_{\text{blood}} \mu_{\text{blood}} + (1-f_{\text{blood}}) \mu_{\text{tissue}} \).

We present results below for fitting \( g_{1,\text{MC}}(\tau) \). To establish the quantitative relationship between \( BF_{\text{conv}} \) and \( BF_{\text{abs}} \) for convective and diffusive RBC displacements independently, we obtain fitting results alternately setting \( D = 0 \) and \( f_{\text{RBC}} = 0 \), respectively. We will distinguish these temporal field autocorrelation functions as \( g_{1,\text{conv}}^\text{max}(\tau) \) and \( g_{1,\text{diff}}^\text{max}(\tau) \) for convective and diffusive RBC displacements, respectively.

In the case of convective motion only, we expect \( \langle v \rangle \) to be the spatially weighted average of RBC speed within the blood vessel. As the absorption length and photon transport length inside the vessel are generally larger than the vessel diameter, we
expect uniform spatial sampling of the vessel and thus expect \( \langle v \rangle = \frac{v_{\text{max}}}{2} \) for a parabolic laminar flow profile. We thus arrive at an equation for \( BF_{\text{conv}} \)

\[
BF_{\text{conv}} = \frac{1}{\frac{1}{2}} \left( \frac{\mu'_s}{\mu'_i} \right)^{\frac{1}{2}} \left( \frac{h_{\text{space}} \pi R}{\mu'_s} \right) \left( \frac{v_{\text{max}}}{R} \right).
\]

In the case of diffusive motion only, we expect \( \langle D \rangle \) to be the spatially weighted average of the RBC diffusion coefficient within the blood vessel, i.e., \( \langle D \rangle = 4\alpha_{\text{blood}} v_{\text{max}} / 3R \) given a parabolic laminar profile and uniform photon sampling of the blood vessel. We thus arrive at an equation for \( BF_{\text{dif}} \)

\[
BF_{\text{dif}} = \frac{4\alpha_{\text{blood}} \mu'_s}{3} \left( \frac{\mu'_s}{\mu'_i} \right)^{\frac{1}{2}} \pi R h_{\text{space}} v_{\text{max}}.
\]

We define absolute blood flow as the volume of blood transiting through a unit volume of tissue per second, which is given by the flow in a vessel (cross-sectional area times average RBC speed) times the number of vessels per unit volume of tissue, i.e.,

\[
BF_{\text{abs}} = \frac{1}{2} \pi R^2 v_{\text{max}} n_{\text{ves}}.
\]

where the number of vessels per unit volume of tissue in our geometry is

\[
n_{\text{ves}} = \frac{1}{h_{\text{space}}^2}.
\]

We thus see that

\[
BF_{\text{conv}} = \left( \frac{\mu'_s}{\mu'_i} \right)^{\frac{1}{2}} \left( \frac{h_{\text{space}} \pi R}{\mu'_s} \right) BF_{\text{abs}}
\]

and

\[
BF_{\text{dif}} = \frac{8\alpha_{\text{blood}} \mu'_s}{3} \left( \frac{\mu'_s}{\mu'_i} \right)^{\frac{1}{2}} \frac{1}{R} BF_{\text{abs}}.
\]

Thus, the prediction from the CDE is that the blood flow index measured by DCS is proportional to absolute blood flow, but the proportionality is modulated by \( \mu'_s/\mu'_i \) and \( R \). Additionally, the proportionality between \( BF_{\text{conv}} \) and \( BF_{\text{abs}} \) is further modulated by the number density of vessels. Note that we assume an estimate of \( \mu'_s/\mu'_i \) from, for instance, an NIRS measurement or from a multidistance DCS measurement.

Finally, we present results below considering the combination of convective and diffusive RBC displacements to establish which dominates the decay of \( g_{1\text{total}}(\tau) \). Given the overwhelming indication from experimental studies, we expect to observe that the diffusive motion dominates the decay.

### 3 Results

#### 3.1 Impact of Scattering Length Within a Vessel

Table 1 summarizes representative optical properties of blood and extravascular tissue at 800 nm for a blood hematocrit of 40%. We use \( g \) and the scattering coefficient \( \mu_s = 1/l_s \) in our Monte Carlo simulations. For the analytical solution provided by the CDE, we use the reduced scattering coefficient \( \mu'_s = \mu_s(1-g) \).

As a preamble, we explicitly test the prediction offered by the CDE that \( g_{1}(\tau) \) is invariant to changes in \( \mu_s \) and \( g \) provided that \( \mu'_s \) is constant. To this end, we ran Monte Carlo simulations for three different combinations of \( \mu'_s, \mu_s, \) and \( g \) maintaining a constant \( \mu'_s = 1.9 \text{ mm}^{-1} \). Specifically, we use \( \mu'_s = 1.9, 19, \) and 190 \text{ mm}^{-1}, and \( g = 0.9, 0.99, \) and 0.99, respectively. The rest of the optical parameters match the values in Table 1, except that we use \( \mu'_s = 1.0 \text{ mm}^{-1} \) and \( g = 0 \), which are typical values we use in our Monte Carlo simulations for the human head to accelerate the computation.

We show results for 80-\text{um}-diameter vessels, spaced every 300 \text{ mm}, with \( v_{\text{RBC,max}} = 2 \text{ mm/s} \).

The results for \( g_{1\text{conv}}(\tau) \) and \( g_{1\text{dif}}(\tau) \) are shown in Fig. 3. These results confirm the prediction from the CDE that, at least for early decay times, \( g_{1}(\tau) \) is invariant as long as \( \mu'_s / (1-g) \) remains constant. The behavior at long decay times does exhibit some dependence on \( \mu_s \) of the blood [see enlargement in panel b].

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**Table 1** Optical properties for blood and extravascular tissue at 800 nm.

<table>
<thead>
<tr>
<th>Property</th>
<th>( \mu_s ) (mm(^{-1}))</th>
<th>( I_s ) (mm)</th>
<th>( \mu_a ) (mm(^{-1}))</th>
<th>( I_a ) (mm)</th>
<th>( g )</th>
<th>( \mu_s' ) (mm(^{-1}))</th>
</tr>
</thead>
<tbody>
<tr>
<td>Blood</td>
<td>0.3</td>
<td>3.3</td>
<td>82</td>
<td>0.012</td>
<td>0.977</td>
<td>1.9</td>
</tr>
<tr>
<td>Tissue</td>
<td>0.002</td>
<td>500</td>
<td>10</td>
<td>0.1</td>
<td>0.9</td>
<td>1.0</td>
</tr>
</tbody>
</table>

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**Fig. 3** (a) The decay of the temporal field autocorrelation function at a 2-cm source–detector separation is shown for \( \mu'_s = 1.9 \text{ mm}^{-1} \) with \( g = 0, 0.9, \) and 0.99. We used 80-\text{um}-diameter vessels, spaced every 300 \text{ mm}, with \( v_{\text{RBC,max}} = 2 \text{ mm/s} \). The dotted, dashed, and solid lines indicate \( g_{1\text{conv}}(\tau) \), \( g_{1\text{dif}}(\tau) \), and \( g_{1}(\tau) \), respectively. (b) An enlargement of the \( 10^{-5} < \tau < 2 \times 10^{-4} \) range in (a).
Table 2. Vascular parameter ranges explored in this study. Each parameter was varied separately while the others were held at a fixed value.

<table>
<thead>
<tr>
<th>Vascular parameter</th>
<th>Range</th>
</tr>
</thead>
<tbody>
<tr>
<td>RBC speed ($v_{RBC}$)</td>
<td>1, 2, 4 mm/s</td>
</tr>
<tr>
<td>Vessel radius ($R$)</td>
<td>30, 40, 50, 60, 70, 80, 90, 100 μm</td>
</tr>
<tr>
<td>Vessel spacing ($h_{space}$)</td>
<td>200, 300 μm</td>
</tr>
<tr>
<td>Hematocrit (Hct)</td>
<td>32%, 36%, 40%, 44%, 48%</td>
</tr>
</tbody>
</table>

Like as $μ_t$ of the blood is decreased, the probability increases that a photon traverses a vessel without scattering from a moving RBC. Importantly, we observe that $g_i^{\text{dif}}(τ)$ decays more quickly than $g_i^{\text{conv}}(τ)$, indicating that the shear-induced diffusion of the RBCs produces more dynamic fluctuations in the electromagnetic field than the flow speed convective displacements. Based on these results, for the remainder of the paper, we use $g_{\text{blood}} = 0.9$ since this gives us effectively the same results we would obtain with $g_{\text{blood}} = 0.977$ and leads to a more efficient execution of the Monte Carlo simulations.

3.2 Blood Flow Indices Estimated from Fitting Monte Carlo Data Using the Correlation Diffusion Equation

In this section, we present results to validate the relations for the blood flow indices $BF_i^{\text{conv}}$ and $BF_i^{\text{dif}}$ given by Eqs. (10) and (11), respectively. We do so by fitting the solution of the CDE, given by Eq. (1), to $g_{i,MC}(τ)$, obtained from the Monte Carlo simulations. As detailed in Table 2, we varied RBC speed, vessel radius, vessel spacing, and hematocrit in these simulations to fully test the dependencies of $BF_i^{\text{conv}}$ and $BF_i^{\text{dif}}$ on each parameter. Note that we fit $g_i^{\text{conv}}(τ)$ to estimate $BF_i^{\text{conv}}$ and we fit $g_i^{\text{dif}}(τ)$ to estimate $BF_i^{\text{dif}}$. We only fit $g_i^{\text{tot}}(τ)$ with the diffusion term because our simulations indicate that the $τ^2$ dependence arising from the convective displacement of the RBCs has negligible impact on $g_i^{\text{tot}}(τ)$ for our parameters of interest. For all cases presented in this section, the blood optical properties were $μ_i = 1.9 \text{ mm}^{-1}$, $g = 0.9$, and $μ_a = 0.30 \text{ mm}^{-1}$. For the tissue, we set $μ_i = 1.0 \text{ mm}^{-1}$, $g = 0.0$, and $μ_a = 0.002 \text{ mm}^{-1}$.

We first present results increasing the speed $v_{\text{max}}$ of the RBCs. Figures 4(a) and 4(b) show $g_{i,MC}(τ)$ and the CDE fit for $v_{\text{max}} = 1, 2,$ and $4 \text{ mm/s}$ for $g_i^{\text{tot}}(τ)$ and $g_i^{\text{conv}}(τ)$, respectively. In this example, the vessel diameter is set to $80 \μm$, the vessels are spaced every $300 \μm$ and the hematocrit is 40%.
set to 40%. These results indicate that the CDE fits $g_{\text{tot}}(\tau)$ and $g_{\text{conv}}(\tau)$ well. Figures 4(c) and 4(d) show the fitted values of $BF_{\text{dif}}$ and $BF_{\text{conv}}$ versus speed, compared with those predicted using Eqs. (10) and (11). These results confirm that the $BF_i$s increase linearly with RBC speed and confirm the accuracy of Eqs. (10) and (11).

Figure 5 shows the $BF$ fitting results for variations in the absolute volumetric blood flow achieved by varying the vessel diameter at a constant maximum RBC speed $v_{\text{max}}$. The vessel diameter was varied from 30 to 100 $\mu$m and results are shown for vessel spacings of 200 and 300 $\mu$m, respectively. As with the previous results, we fit $g_{\text{tot}}(\tau)$ and $g_{\text{conv}}(\tau)$ to estimate $BF_{\text{tot}}$ and $BF_{\text{conv}}$, respectively. We then used Eqs. (14) and (15) to determine the expected $BF_i$ values and find that this predicts well the measured $BF_i$.

In Fig. 6, we plot fitted $BF_i$ values for changing the blood hematocrit by $\pm 20\%$ from our baseline value of 40%. In this range, the reduced scattering coefficient of blood is changing linearly with hematocrit. We also show the expected $BF_i$ values and find excellent agreement.

4 Discussion

The agreement between the prediction by the CDE and the fits to the Monte Carlo results is striking. Despite the heterogeneous optical and dynamic properties of the vascular structure and the correlated consecutive scattering events within a vessel, the homogeneous solution of the CDE accurately describes the decay of $g_i(\tau)$. We have shown results fitting $g_{\text{tot},\text{MC}}(\tau)$ and $g_{\text{conv},\text{MC}}(\tau)$, allowing us to test the convective and diffusive RBC displacement models for the CDE and demonstrate that the diffusive motion of RBCs dominates the decay of $g_{\text{tot},\text{MC}}(\tau)$. Our fitting results confirm that Eqs. (14) and (15), which relate $BF_{\text{conv}}$ and $BF_{\text{dif}}$ to blood flow, are accurate within 5%. Importantly,
while the CDE allows us to write a linear relation between the estimated blood flow index and true absolute blood flow, we see that the proportionality is modulated by the radius of the blood vessels and the reduced scattering coefficient of the blood (which is modulated by hematocrit). In the case of BF_{if}, the proportionality is further dependent on the number density of vessels. In this sense, it is fortunate that the diffusive motion of RBCs dominates the decay of g_{MC}(τ); thus, we do not need knowledge of the number density of vessels to estimate BF_{abs}. Recall that the BF_{S} depend on hematocrit even when absolute blood flow is constant because the BF_{S} depend on the probability of scattering from an RBC, Eq. (9), which increases linearly with the reduced scattering coefficient of blood.

Given that the diffusive motion of RBCs dominates the decay of g_{MC}^{s}(τ), Eq. (15) thus stands as the quantitative relationship between the blood flow index measured by DCS and the true absolute blood flow. Recall that in order to get BF_{S}, we first need to know the average optical properties of the tissue \(\mu_{a,avg}\) and \(\mu_{s,avg}\), which would generally be obtained from an independent NIRS measurement\(^{31,32}\) and possibly also estimated with multidistance DCS measurements.\(^{30}\) To then convert the BF_{if} to BF_{abs}, we need to know the proportionality \(\alpha_{shear}\) between shear flow and the RBC diffusion coefficient, the reduced scattering coefficient of the blood \(\mu_{s,blood}\), and the radius of the blood vessels. In this paper, we used a value of \(\alpha_{shear} = 10^{-6} \text{ m}^2\text{s}^{-1}\) obtained from Goldsmith and Marlow.\(^{15}\) We note that this is a value obtained for a hematocrit of 40% to 47% and that \(\alpha_{shear}\) was found to linearly increase with hematocrit from 0% to 45% and then to plateau and reverse.\(^{37}\) The reduced scattering coefficient of blood \(\mu_{s,blood}\) is linearly proportional to hematocrit from 0% to 40%, the highest value measured by Meinke et al.\(^{21}\) In principle, if one has an independent measure of hematocrit, from a blood draw for instance, one can then determine the appropriate value to use for \(\alpha_{shear}\) and \(\mu_{s,blood}\). The remaining factor needed to estimate BF_{abs}, is the vessel radius \(R\). Although we performed simulations using vessels with a common radius, in reality DCS will measure vessels with a distribution of radii. We anticipate that Eq. (15) will still be valid when measuring a distribution of vessel radii, but that the effective vessel radius will represent a complex nonlinear dependence on the distribution of vessel radii and the corresponding RBC speed distribution.

We note that in our previous publication,\(^{13}\) we postulated that the diffusional dependence of the DCS measurements in tissue arose because of sequential multiple scattering within a single vessel. As demonstrated in this paper, this factor does not play a major role in the interplay between diffusive and convective influences on the autocorrelation of multiple scattered light. Rather, it is simply that at source–detector separations commonly used for DCS measurements, the shear-induced diffusion dominates the signal. We estimate that the convective contribution only becomes significant when the detected photon path lengths become less than a millimeter, but this requires further investigation. A potential implication is that the convective contribution should be considered in analyzing laser speckle contrast imaging data.\(^{34}\) It is worth noting at this point that the concept of a well-defined absolute blood flow has been questioned as the value of absolute blood flow obtained has been theoretically demonstrated to depend on vascular geometry, measurement procedure, and spatial scale sampled.\(^{35,36}\) Further research is needed to better understand the implications of these theoretical arguments.

5 Summary
We have presented a set of Monte Carlo simulations that quantify the contribution of diffusive and convective RBC motion to the decay of the autocorrelation function of light that has scattered through tissue. Through this modeling, we demonstrate that diffusive motion dominates the autocorrelation decay for typical experimental parameters used in DCS measurements. We have also provided expressions for the expected dependence of the measured DCS blood flow index on absolute tissue blood flow in terms of the vascular volume fraction, average vessel diameter, and blood optical properties. Our results indicate that the DCS blood flow index commonly obtained using the established CDE indeed offers a direct measure of tissue blood flow, but the proportionality is also sensitive to changes in hematocrit and in the average vessel diameter in the region probed by light. The immediate implication is that care must be taken to control for variations in these additional parameters when reporting DCS blood flow measurements. We expect further studies to explore these sources of variability and help define the range of effective DCS applications.

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