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Effects of Flowing RBCs on Adhesion of a Circulating Tumor Cell in

Microvessels

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Abstract

Adhesion of circulating tumor cells (CTCs) to the microvessel wall largely depends on the blood hydrodynamic conditions, one of which is the blood viscosity. Since blood is a non-Newtonian fluid, whose viscosity increases with hematocrit, especially in microvessels with low shear rates. In this study, the effects of hematocrit, vessel size, flow rate and red blood cells (RBCs) aggregation on adhesion of a CTC in the microvessels were numerically investigated using dissipative particle dynamics. The membrane of cells was represented by a spring-based network connected by elastic springs to characterize its deformation. RBCs aggregation was modelled by a Morse potential function based on depletion-mediated assumption and the adhesion of the CTC to the vessel wall was achieved by the interactions between receptors and ligands at the CTC and those at the endothelial cells forming the vessel wall. The results demonstrated that in the microvessel of 15µm diameter, the CTC has an increasing probability of adhesion with the hematocrit due to a growing wall-directed force, resulting in a larger number of receptor-ligand bonds formed on the cell surface. However, with the increase in microvessel size, an enhanced lift force at higher hematocrit detaches the initial adherent CTC quickly. If the microvessel is comparable to the CTC in diameter, CTC adhesion is independent of the *Hct*. In addition, the velocity of CTC is larger than the average blood velocity in smaller microvessels and the relative velocity of CTC decreases with the increase in microvessel size. An increased blood flow resistance in the presence of CTC was also found. Moreover, it was found that the large deformation induced by high flow rate and the presence of aggregation increase the CTC adhesion.

Keywords: Adhesion · Aggregation · Circulating tumor cell · Dissipative particle dynamics · Red blood cell

1 Introduction

CTCs are caused by the tumor cell intravasation into the blood stream from the original tumor. Such CTCs are transported by the blood flow to the distant organs through the vasculature to form secondary tumors. During the blood circulation, the tumor cell must suffer the hydrodynamic shear stresses, and local forces due to neighboring cells. In order to step out of the flow, a very important step is the so called "adhesion cascade" (Guo et al. 2004; Haier and Nicolson 2001; Stroka and Konstantopoulos 2014; Wirtz et al. 2011). Most previous studies have experimentally (Cheung et al. 2011; Fu et al. 2015; Guo et al. 2014; Yan et al. 2012; Zhang et al. 2016) and numerically (Rejniak 2012; Yan et al. 2012; Yan et al. 2010) investigated the process of cell adhesion to the blood vessel. The numerical studies simply regarded the blood as a homogenous Newtonian fluid (Rejniak 2012; Yan et al. 2010). However, under shear flow, shear stresses acting on the CTCs largely depend on blood viscosity. As a matter of fact blood is a non-Newtonian fluid and its viscosity increases markedly with the increase of hematocrit at low shear rates. In addition, CTCs rolling along the vessel wall can greatly increase the vascular resistance and modify the blood flow and biophysical conditions to affect adhesive interactions. Although the interactions between the tumor cell and red blood cells in the blood stream remain unclear, some similarities with the interactions between white blood cells (WBCs) and RBCs may be found.

The moving RBCs were proved to promote WBC adhesion either by causing margination of leukocytes or by initiating adhesion and stabilizing attachment (Abbitt and Nash 2003; Munn et al. 1996). During microcirculation, the tendency of RBCs flowing to the center of the vessels tends to push leukocytes toward the endothelium, leading to the process of margination. Previous studies have shown that leukocyte margination largely depends on the blood properties, including shear rate, hematocrit and RBC aggregation. Fedosov and Gompper (2014) numerically analyzed the mechanism of three-dimensional leukocyte margination toward the vessel wall and showed that WBC marginates to the wall at a region of intermediate hematocrits, Hct=0.2~0.4, under relatively low flow rates corresponding to the venule shear rate and RBCs aggregation is unnecessary for margination at high hematocrit values. Likewise, Takeishi et al. (2014) showed that leukocytes rarely approach the wall surface within microvilli length (less than 0.5µm) when numerically investigated margination of leukocytes at arteriole shear rate (670/s), but lower shear rate enables the leukocytes contact the wall. Similarly, shear flow induces the deformation of CTCs and the margination of CTCs towards the vessel wall is also expected primarily in the venular part of microcirculation, which would imply that the tissue invasion by tumor cells present in blood largely occurs from venules (Firrell and Lipowsky 1989). In fact, CTCs prefer to adhere to the small venules with relatively low blood flow rates (Guo et al. 2014; Zhang et al. 2016). Moreover, the increase of WBC margination with reduction in shear rate (<50s⁻¹) (Nash et al. 2008) is strongly dependent on the occurrence of RBC aggregation. Also increasing the extent of RBC aggregation also increased the firm adhesion of WBCs to the endothelium because of an enhanced probability of contact between leukocytes and the vessel wall (Abbitt and Nash 2003; Munn and Dupin 2008; Pearson and Lipowsky 2000; Sun and Munn 2006). In addition to this, the leukocyte adhesion increases at the hematocrit ranging from 10% to 30% (Abbitt and Nash 2003). It has been found that the WBC rolling speed increases with RBC concentration because it is affected by the RBC-WBC collision frequency and blunting of the velocity profile (Sun and Munn 2005). Still, the rolling velocity of a WBC is strongly dependent on the local leukocyte WBC concentration. The velocity fluctuation decreases as the WBC concentration increases. Equally, the rolling velocity and velocity fluctuation decreases with increasing cell deformability (Pappu et al. 2008). Recently, the forces exerted by

the flowing blood on an adherent rigid WBC was estimated and it is found that RBCs not only enlarge the streamwise forces, impairing the WBC binding, but also cause an average wall-directed force, which is expected to enhance binding (Isfahani and Freund 2012).

Based on many similarities in the process of margination and adhesion, models developed for leukocytes can be applied to CTCs. King et al. (2015) experimentally and numerically investigated the CTC dynamics in the microvasculature and the numerical findings demonstrated that single CTC with more rigid membrane marginates quicker than the softer one while the experimental results showed that the rolling velocity of cancer cells increases with the number of cells or aggregates. However, for the process of adhesion, softer cell flattening enables adherent cells to avoid high shear forces within the center of the parabolic curve of fluid flow. Takeishi et al. (2015) demonstrated that the cell margination depends on cell relative size to the vessels. When the discrepancy in the radiuses between vessel and tumor cell exceeds the thickness of red blood cell, tumor cell tends to margination. In addition to tumor cell properties, the vessel geometry could contribute to tumor cell adhesion (Weiss 1992). In vivo experiments (Guo et al. 2014; Yan et al. 2012; Zhang et al. 2016) have demonstrated that tumor cells prefer to adhere to the curved vessels and at the bifurcations of microvasculature. Previous findings provide valuable premises for the tumor cell firm adhesion. As limited studies have been conducted on how the tumor cell switches from the initial attachment to firm adhesion under the forces of RBCs, we focus on the effects of flowing RBCs on the adhesion of a CTC to the vessel wall.

In this study, the blood was modeled by a suspension of RBCs. Dissipative particle dynamics method combined with a spring-based network cell model was employed to carry out 3D simulations of motion of a rolling tumor cell in microvessels constructed by the straight tubes. Cell-cell interaction was represented by a Morse-potential function based on a depletion model. Then the simulation results of motion of tumor cell

in blood flow at different hematocrits in the tube ranging from 10 to 20µm were given. Next, the effects of hematocrit, vessel size, tumor cell deformability as well as RBC aggregation on the motion of the rolling circulating tumor cell were investigated. Finally, summary and conclusions were presented.

2 Model and methods

The blood flow is modeled by using dissipative particle dynamics, which has been adopted in our previous study (Xiao et al. 2016). Details of DPD method have been introduced (Espanol 1995; Groot and Warren 1997; Hoogerbrugge and Koelman 1992). In brief, each DPD particle represents a soft lump of atoms and interacts with surrounding particles through three simple pairwise additive forces: conservative force, dissipative force and random force. Particle motion follows Newton's law. In the following, the cell models employed in the simulations are specifically described.

2.1 Cell membrane model

In simulations, the cell membrane is discretized into a collection of particles connected by elastic springs. A spring-based network model endowed with in-plane and bending energy as well as constraint of surface area and volume has been introduced by Boey et al. (1998) to describe RBC initially. A systematic coarse-grained procedure was introduced by Pivkin and Karniadakis (2008) to reduce the number of degrees of freedom dramatically in the RBC model. This coarse-grained model was further improved by Fedosov et al. (2010), yielding accurate mechanical response. The total energy of the network is defined as

$$E(\{\mathbf{r}_i\}) = E_{in-plane} + E_{bending} + E_{area} + E_{volume}$$
(1)

where \mathbf{r}_i represents the vertex coordinates and the in-plane elastic energy for WLC-POW model is given by

$$E_{in-plane} = \sum_{all \, edges} \left[\frac{k_B T l_{max}}{4p} \frac{3x_l^2 - 2x_l^3}{1 - x_l} + \frac{k_p}{(m-1)l^{m-1}} \right]$$
(2)

where $x_l = l/l_{max} \in (0,1)$, l_{max} is the maximum spring extension, which is equal to 2.2 times equilibrium spring length for the WLC model, p is the persistence length, k_B is Boltzmann constant and T is temperature of the system, which is equal to 310K. k_p is a POW force coefficient and m is a specified exponent, here we set it to 2 (Fedosov et al. 2010).

The bending energy stored in the adjacent triangular elements is defined by

$$E_{bending} = \sum_{all \ triangle \ adjacents} k_{bend} \left[1 - \cos(\theta_{\alpha\beta} - \theta_0) \right]$$
(3)

where k_{bend} is a bending modulus; $\theta_{\alpha\beta}$ is the instantaneous angle formed between the outer normal vectors of two adjacent triangles α , β sharing the common edge; θ_0 is the spontaneous angle.

The area and volume conservation constraints are

$$E_{area} = \frac{k_{area}^{tot} (A^{tot} - A_0^{tot})^2}{2A_0^{tot}} + \sum_{all \, triangles} \frac{k_{area} (A - A_0)^2}{2A_0}$$
(4)

$$E_{volume} = \frac{k_{volume} (V - V_0^{tot})^2}{2V_0^{tot}}$$
(5)

where k_{area}^{tot} , k_{area} and k_{volume} are constraint constants for global area, local area, and volume; A^{tot} and V are the instantaneous membrane area and the cell volume; A_0^{tot} and V_0^{tot} are their respective specified total area and volume values. A, A_0 are the instantaneous and initial local area.

Nodal forces are derived from the total energy as follows:

$$\mathbf{F}_{i}^{\text{membrane}} = -\partial \mathbf{E}\{(\mathbf{r}_{i})\} / \partial \mathbf{r}_{i}$$
(6)

The elasticity of the network is based on the linear analysis of a two-dimensional sheet of springs built with equilateral triangles (Dao et al. 2006). The linear shear modulus of the WLC-POW model is

$$G = \frac{\sqrt{3}k_BT}{4pl_{\max}x_0} \left(\frac{x_0}{2(1-x_0)^3} - \frac{1}{4(1-x_0)^2} + \frac{1}{4}\right) + \frac{\sqrt{3}k_p(m+1)}{4l_0^{m+1}}, \quad x_0 = l_0 / l_{\max}$$
(7)

The linear area compression modulus is defined as

$$K = 2G + k_{area}^{tot} + k_{area} \tag{8}$$

The Young's modulus *Y* for the two-dimensional sheet can be expressed through the shear and areacompression moduli as follows

$$Y = \frac{4KG}{K+G} \tag{9}$$

And the Poisson's ratio v is given by

$$\nu = \frac{K - G}{K + G} \tag{10}$$

Based on the incompressibility assumption $k_{area}^{tot} + k_{area} >> G$ is set, so $Y \to 4G$ and $v \to 1$.

The relationship between bending modulus k_{bend} and the macroscopic membrane bending rigidity k_c is derived for the case of a spherical shell in the Helfrich bending energy, as follows:

$$k_{bend} = \frac{2}{\sqrt{3}} k_c \tag{11}$$

2.2 RBC aggregation model

To analyze the RBC aggregation, the depletion model introduced by Liu et al. (2004) was employed. In this model a Morse potential function $\psi(r)$ was used to model the interaction energy

$$\psi(r) = D_e[e^{2\beta(r_0 - r)} - 2e^{\beta(r_0 - r)}]$$
(12)

where *r* is the distance between two plane elements of the opposing RBCs directly facing each other, r_0 is the zero force length, D_e is the intercellular interaction strength, and β is the scaling factor controlling the interaction decay behavior. Therefore, the total interaction energy (Ye et al. 2014) of a triangulated cell is expressed by

$$\psi(\{\mathbf{r}_i\}) = \sum_{j=1,N_t} \psi(r_{jk}) (\mathbf{n}_j \cdot \mathbf{I}_j) (\mathbf{n}_k \cdot \mathbf{I}_k) A_j$$
(13)

where r_{jk} is the local distance between the *j*th and kth triangles located in cells 1 and 2 respectively. N_i is the number of the triangle elements which are linked by particle *i*. \mathbf{n}_j and \mathbf{n}_k are the outward unit normal vectors to the those curved elements, \mathbf{I}_j and \mathbf{I}_k stand for the unit vectors parallel to the line joining the centers of two cells and directed toward each other, which are based on the DLVO (Derjaguin-LandauVerwey-Overbeek) theory (Bhattacharjee et al. 1998) to describe the interaction energy between two curved surfaces. A_j is the area of *j*th triangle of cell 1. The interaction force acting on the membrane particle *i* in cell1 is given by:

$$\mathbf{F}_{i}^{\text{agg}} = -\frac{\partial \psi(\{\mathbf{r}_{i}\})}{\partial \mathbf{r}_{i}}$$
(14)

It has been shown that the intercellular force is simply illustrated as a weak attractive force at far distance, but a strong repulsive force at near distance. In the numerical model, in order to save the computational cost, a cut-off distance r_c^{agg} is defined as zero force separation threshold, set to be 1.5, beyond which the intercellular force is assumed to be zero.

2.3 CTC adhesion model

Adhesion of cells to the vessel wall is mediated by the interactions between receptors and ligands at the CTC and those at the endothelial cells forming the vessel wall. The adhesion model provides the rules of formation and dissociation of bonds between receptors and ligands. The probabilistic model developed by Hammer and Apte (1992) is commonly used in simulations and is known as adhesive dynamics. If the distance between a receptor and a free ligand is less than the reactive distance d_{on} , a new bond may form with the association rate k_{on} . A free ligand refers to that it is not bound to any receptors. A preexisting bond is ruptured with off-rate k_{off} or if its length exceeds the rupture distance d_{off} . The rates k_{on} and k_{off} are computed as:

$$k_{on} = k_{on}^{0} \exp(-\frac{\sigma_{on}(l^{b} - l_{0}^{b})^{2}}{2k_{B}T}),$$

$$k_{off} = k_{off}^{0} \exp(\frac{\sigma_{off}(l^{b} - l_{0}^{b})^{2}}{2k_{B}T})$$
(15)

where k_{on}^0 and k_{off}^0 are the unstressed reaction rates at the distance $l^b = l_0^b$ between a receptor and a ligand with the equilibrium spring length l_0^b . The effective on and off strengths σ_{on} and σ_{off} define a decrease or an increase of the corresponding rates within the interaction lengths d_{on} and d_{off} . A spring

force acted on the receptors and ligands linked by an existing bond is calculated from the Hooke's law

$$F(l^{b}) = k_{s}(l^{b} - l_{0}^{b})$$
(16)

where k_s is the spring constant. The binding force can be derived as $\mathbf{F}_i^{\text{bond}} = F(r_{ij})\hat{\mathbf{r}}_{ij}$. The probability of the formation of a new bond and that of the breakage of an existing bond are given by

$$P_{on} = \begin{cases} 1 - e^{-k_{on}\Delta t}, l^{b} < d_{on}, \\ 0, l^{b} \ge d_{on}, \end{cases} \qquad P_{off} = \begin{cases} 1 - e^{-k_{off}\Delta t}, l^{b} < d_{off} \\ 0, l^{b} \ge d_{off} \end{cases}$$
(17)

where Δt is the time interval. At a given time instance, two random numbers ξ_1 and ξ_2 uniformly distributed on [0, 1] are generated. An existing bond is ruptured if $\xi_1 < P_{off}$ and left unchanged otherwise. A bond formation is allowed for each found receptor according to the relationship $P_{on} > \xi_2$.

2.4 Model and physical units scaling

In order to keep the simulation system consistent with the real system, the physical properties should be mapped onto the dimensionless properties in the model. A scaling procedure has been presented by Fedosov et al. (2010), which relates the model's non-dimensional units to physical units. The length scale is given by:

$$L^{S} = \frac{D_0^P}{D_0^M} \tag{18}$$

where the superscripts M and P denote "model" and "physical". The energy scale is provided as follows

$$E^{S} = \frac{Y^{P}}{Y^{M}} \left(\frac{D_{0}^{P}}{D_{0}^{M}}\right)^{2}$$
(19)

The force scale is defined by

$$N^{S} = \frac{Y^{P}}{Y^{M}} \frac{D_{0}^{P}}{D_{0}^{M}}$$
(20)

The scaling between model and physical times is defined as follows

$$\tau^{S} = \frac{Y^{M}}{Y^{P}} \frac{D_{0}^{P}}{D_{0}^{M}} \frac{\eta^{P}}{\eta^{M}}$$
(21)

3 Results and discussion

3.1 Simulation setup and modeling parameters

The tumor cells arrest and adhesion experiment was performed, in which the individual trajectories of tumor cells in the blood flow can be measured by labeling them with the fluorescent dye. A certain portion of tumor cells were found to be arrested at the postcapillary venules, as shown in Fig. 1, where there is no size restriction and the flow rate is relatively low. Based on the experimental observation, the motion of a rolling tumor cell under blood flow with a low velocity in a microvessel with a diameter ranging from $D = 10 \sim 20 \mu \text{m}$ and a length of $L = 45 \mu \text{m}$ were simulated, as illustrated in Fig. 2. The blood was modeled as a suspension of RBCs. Each RBC consists of 640 particles and is 7.82 µm in diameter. It has a thickness at the thickest point of 2.5µm and a minimum thickness in the center of 1µm. The tumor cell is of spherical shape with a diameter of 9 µm, constructed by 1170 particles. The nucleus of tumor cell is neglected for simplicity in this study. Both the RBCs and CTC are filled with a Newtonian fluid with a viscosity of $\mu_i = 6 \times 10^{-3} Pa \cdot s$.



Fig. 1 CTC adhesion at a postcapillary venule. The adherent CTC is labeled with the fluorescent dye. The blood flow

direction is indicated by a yellow arrow.



Fig. 2 Schematic illustration of computational domain

The parameters of plasma, RBCs and CTC are listed in Table 1. No-slip boundary condition was imposed near the wall and a periodic boundary condition was applied along the flow direction. Bounce-back reflection was exerted on the surface of the cell membrane to ensure its impenetrability. To drive the flow, a uniform body force was applied to all particles in the flow direction, which is equivalent to the pressure gradient $\Delta P/L = \rho f$, where ΔP is the pressure drop over the tube length and ρ is the suspension's mass density. As the tumor cell margination and adhesion is expected to occur at a lower flow rate, the shear rates are set to below 20s⁻¹.

Tuble Tomanulon parameters for con monitrate and plasma							
Parameter	Simulation	Physical values					
Blood plasma density ($ ho$)	6	1.0×10^3 kg / m ³ (Skalak and Jian 1987)					
Blood plasma viscosity (μ)	20.4	1.2×10^{-3} Pa · s (Skalak and Jian 1987)					
Temperature (T)	e (T) 0.0828 310K						
Membrane Young's modulus for RBCs (Y^{R})	369	$18.9 \mu N/m$ (Suresh et al. 2005)					
Membrane Young's modulus for RBCs (Y^T)	80.5~8050	$4.16 \sim 416 \mu N/m$					
Membrane bending modulus for RBC (k_{bend}^R)	5.364	2.8×10^{-19} J (Fedosov et al. 2010)					
Membrane bending modulus for RBC (k_{bend}^T)	69.28	$3.6 \times 10^{-18} \mathrm{J}$					
Intercellular energy density (D_e)	3.872	$0.2\mu J/m^2$ (Zhang et al. 2009)					
Scaling factor (eta)	7.68	$7.68 \mu m^{-1}$					
Zero force distance (r_0)	0.3	0.3μm (Fedosov et al. 2011)					
Time step (Δt)	0.001	0.00114ms					
Global area constraint constant (k_{area}^{tot})	5×10^{4}	$2.58 \times 10^{-3} \text{N/m}$					
Local area constraint constant (k_{area})	100	5.2µN/m					
Volume constraint constant (k_{volume})	5×10^{4}	$3.35 \times 10^{-3} \text{N/m}$					
Unstressed on rate (k_{on}^0)	11.3	10 ⁴ s ⁻¹ (Schwarz and Alon 2004)					
Unstressed off rate (k_{off}^0)	0.023	20 s ⁻¹ (Alon et al. 1997)					
On strength (σ_{au})	9.68	0.5µN/m (Dembo et al. 1988)					

Table 1 Simulation parameters for cell membrane and plasma

Off strength (σ_{off})	0.968 0.05µN/m (Dembo e		
Association length (d_{on})	0.1	0.1µm	
Disassociation length(d_{off})	0.1	0.1µm (Marshall et al. 2006)	
Spring constant (k_s)	1.55×10^{5}	8×10^{-3} N/m (Fritz et al. 1998)	
Equilibrium spring length(l_0^b)	0.025	0.025µm (Dembo et al. 1988)	
Receptor density (n_r)	4.6	4.6/ μm ²	
Ligand density (n_l)	4	4 /µm ²	

Initially, CTC was released near the wall with a distance of 50nm to facilitate binding by the receptorligand formation. When the flow was turned on, the suspension began to flow and the adherent tumor cell rolled along the wall. Representative snapshots of the simulated CTC and RBCs motion within the microvessels with a diameter of 15µm were presented in Fig. 3. The RBCs are scattered in the flow at low flow rates, so the particulate nature of the blood is pronounced. The CTC was found to move faster than the RBCs by comparing the trajectories of CTC and the green RBC. When the RBCs were overtaken by the CTC, most RBCs approached the upper end of the CTC. The forces induced by the collision of these RBCs push the CTC toward the wall in further, as illustrated in Fig. 3d.



Fig. 3 Representative images of the motion of a CTC (blue) and RBCs (red) in the microvessel with a diameter of 15µm

at (a) t=0s; (b) t=0.25s; (c) t=0.5s; (d) t=0.75s. The green RBC represents a reference

As the adhesion of CTC in the blood flow is a complex process, which is greatly affected by the

properties of the blood, the following sections will analyze the effect of hematocrit, CTC deformability, vessel size, flow rate as well as RBCs aggregation on the CTC adhesion in detail.

3.2 Dependence on hematocrits and tumor cell deformability

The motion of a rolling CTC in the flow with hematocrits ranging from 0.1 to 0.3 in a microvessel of 15μ m in diameter was simulated firstly. The results demonstrate that at a lower hematocrit (*Hct* = 0.1, 0.2), the rolling tumor cell might be pulled off the vessel by the intercellular interactions, as shown in Fig. 4a. However, at *Hct* = 0.3, the CTC maintains to roll along the wall (Fig. 4b), resulting in a larger number of receptor-ligand bonds formed on the cell surface compared with the cases of *Hct* =0.1 and *Hct* =0.2, as shown in Fig. 4c. The force on the CTC is defined by the cumulative force generated by the interactions with the RBC and plasma. The positive force implies a lift force pointing to the vessel center while the negative value refers to a wall-directed force. Due to the particulate nature of blood, the CTC has a lower collision frequency with RBCs and subjects to a weaker wall-directed force (Fig. 4d) at lower hematocrits. However, at *Hct* = 0.3, a stronger wall-directed force inhibits detachment of CTC from the vessel wall. The maximum pressure normal to the wall on the wall-leukocyte has been estimated as about 30Pa (Isfahani and Freund 2012). This pressure is equivalent to the maximal force divided by the CTC surface area, so the value of 118Pa was obtained, which is larger than that estimated value. That is because the CTC is larger and the hematocrit is higher compared to the simulations of *Hct* =0.25 in (Isfahani and Freund 2012).

(a)



(b)



(c)



Fig. 4 Simulation snapshots of RBCs (red) and a circulating tumor cell (blue) at Hct=0.2 (a) and Hct=0.3(b); and the variations of number of receptor-ligand bonds formed on the surface of CTC (c) and of the force on the CTC (d) at different hematocrits in a microvessel of 15 μ m in diameter

Then, the effects of CTC deformability on its deformation and adhesion in a microvessel of 15µm diameter were investigated. Three values of elastic modulus $Y = 416\mu$ N/m, 41.6μ N/m and 4.16μ N/m were employed to represent different deformabilities. To quantify the deformation of a rolling CTC, the deformation indexes H_T / D_T and L_T / D_T to characterize the CTC deformation along the radial direction and flow direction respectively, are shown in Fig. 5a and Fig. 5b. It can be found that there is a slighter fluctuation in the evolution of the deformation in the CTC with $Y = 416\mu$ N/m. But when the Young's modulus decreases by 10 times, the CTC deforms significantly within 8%. The deformation during rolling may be of importance in adhesion owing to their direct effect on contact area correlated with the number of receptorligand bonds. Figure 5c shows the time history of number of receptor-ligand bonds formed on the surface of the CTC with different deformabilities. It seems that the softer cell with $Y = 41.6 \mu N/m$ has the largest number of bonds compared to the other two cases especially after t=0.2s. This is because that the deformation of this soft cell along tube axis is larger than the stiff one ($Y = 416\mu$ N/m). For the softest cell, the lift force increases the cell height, leading to the decline in the length and the number of bonds. Despite the discrepancies on the deformation for the CTC with different deformabilities have been observed, the deformation is so small by virtue of the low shear rate, compared to the deformation up to 40% at 800s⁻

¹(Pearson and Lipowsky 2000). Consequently, the deformability has no significant effect on the CTC displacement along the flow direction, as shown in Fig. 5d. The CTC was found to move at a constant velocity.



Fig. 5 Deformation of a rolling CTC (a) along the radial direction H_T/D_T and (b) along the flow direction L_T/D_T , and the receptor-ligand number of bonds formed on the cell surface (c) as well as cell position in the flow direction (d) at Hct =0.3 in a microvessel of 15 µm as a function of CTC membrane deformability. H_T , L_T and D_T denote cell height, length, and initial diameter respectively.

3.3 Dependence on the microvessel size

Since the vessel size has a great effect on the CTC adhesion in blood flow (Takeishi et al. 2015; Wirtz et al. 2011), the flow of the CTC in different microvessels of various diameters ranging from 10 to 20µm were simulated. The blood flow parameters are summarized in Table 2. The mean flow velocity is $V_{average} = Q/A = \int_{A} v(r) dA_c / A_c$, where Q is volume flow rate and A_c is the area of cross section. The mean shear rate is represented by $\gamma = v_{average}/D$, and pressure gradient is expressed by $\Delta P/L$. As the CTC adhesion

is expected at lower shear rates, the mean shear rates are less than $20s^{-1}$. The capillary number defined by $Ca = \mu_i \gamma D^R / 2G^R$, is less than 0.075. Here, R represents the RBC, the diameter of RBC is $D^R = 7.82 \mu m$, the shear modulus for RBC is $G^R = 6.3 \mu N/m$.

Table 2 Blood flow parameters for different hematocrit values and different tube diameters. *D* is the tube diameter, *Hct* is the tube hematocrit, γ is the mean shear rate, $\Delta P/L$ is the pressure gradient, v_p is the average velocity for the blood flow

without CTC								
<i>D</i> (µm)	Hct	v _{average} (mm/s)	γ (1/s)	$\Delta P/L$ (Pa/m)	$v_p (\text{mm/s})$			
10	0.1	0.116	11.63	2.44E+05	0.140			
10	0.2	0.074	7.43	2.44E+05	0.137			
10	0.3	0.043	4.31	2.44E+05	0.121			
15	0.1	0.131	8.74	1.08E+05	0.139			
15	0.2	0.076	5.04	1.08E+05	0.099			
15	0.3	0.040	2.66	1.08E+05	0.054			
20	0.1	0.121	6.05	6.10E+04	0.084			
20	0.2	0.070	3.51	6.10E+04	0.071			
20	0.3	0.044	2.20	6.10E+04	0.051			

For $D = 20\mu m$, the initial adherent CTC is more likely to detach from the vessel at higher hematocrit, which can be seen in Fig. 6a. Figure 6c shows the variation of number of bonds at different hematocrits for $D = 20 \ \mu m$. The initial adherent CTC can be found to detach from the vessel for all cases during rolling along the wall. But at higher hematocrit, the CTC cannot arrive at the vessel again during this simulation time period. Under the low shear flow rate, the weak tendency of RBC flowing towards the center of microvessel inhibits the margination of CTC. Instead, the effect of particulate nature is so pronounced that RBCs in the blood flow are distributed separately and the number of RBCs near the wall is larger compared to the lower hematocrits, leading to an enhanced lift force (Fig. 6d). Therefore, the CTC nearly cannot reach the vessel wall at high hematocrit (*Hct* = 0.3), as shown in Fig. 4a. While at a low concentration of RBCs (*Hct* = 0.1), the detached CTC can easily reapproach the vessel wall (Fig. 6b) under the pushing force exerted by the RBCs. (a)

00

04

t (s)

0 6



Fig. 6 Simulation snapshots of RBCs (red) and a circulating tumor cell (blue) at Hct=0.3 (a) and Hct=0.1 (b); and variation of number of receptor-ligand bonds formed on the surface of CTC (c) and of the force on the CTC (d) at different hematocrits in a microvessel of 20µm in diameter

-6

0.0

0.2

0.6

0.4

t (s)

0.8

To trace the radial position of the CTC in the blood flow, a separation distance δ is introduced, which is defined by the closest distance between the CTC membrane and the vessel wall surface. Here, $\delta \leq 0.1 \mu m$ indicates that the CTC attaches to the vessel wall as the association length is set to 0.1 µm. Figure 7 illustrates the probability of separation distance computed over a time period of 0.9s for D = 10, 15 and 20 μ m at the hematocrits ranging from 0.1 to 0.3. From the bar chart, it can be seen that for $D = 10\mu m$, the initial adherent CTC nearly attaches to the vessel wall, which is independent on the hematocrit. This is due to the fact that the CTC has a stronger confinement $D_T / D = 0.9$. That's why the tumor cell extravasation is expected in the microvessels with a diameter less than the cell diameter. With the decrease in the hematocrit, the probability of CTC detachment from the vessel improves for $D = 15 \mu m$. Interestingly, the relation between the probabilities of separation length and hematocrits for $D = 15 \mu m$ is reverse to that for $D = 20 \mu m$, which is consistent with the findings on the variation of formed bonds number.

(b)



Fig. 7 Probability of the separation distance between the CTC membrane and the wall surface for different sized vessels at Hct = 0.1, 0.2 and 0.3

3.4 CTC velocity and effect of CTC motion on the flow resistance

The velocity of CTC is an important indicator of CTC adhesion especially in larger microvessels. If the CTC maintain attaching the vessel, its velocity may be lower than the average blood velocity. Otherwise, according to the Fahraeus effect, an increases discharge hematocrit is expected when the cells leave the outlet due to its tendency of migrating toward the vessel center, the CTC flows faster than the mean blood. The translational velocity of CTC was examined by normalizing the CTC velocity V_T to the average blood velocity $V_{average}$. Figure 8 shows that the normalized CTC velocity for different values of *Hct* and diameter. It suggests that in smaller microvessels ($D \le 15\mu$ m), the CTC flows faster than the average blood velocity, also can be observed in Fig. 3. This is consistent with the previous finding of $V_T/V_{average} > 1$ when $D/D_T < 2.0$ (Takeishi et al. 2015). That is because that the CTC may be located nearer to the center of the vessel, even attached to the vessel. But when the vessel diameter increases further, the normalized CTC velocity decreases to around the unity at higher hematocrits. Particularly, the CTC moves slower than the blood stream at *Hct* =0.1 in microvessel of 20 μ m, which means that the CTC probably moves near the vessel wall. Moreover, the normalized CTC velocity increases with hematocrit, but this tendency becomes weak in

larger microvessel. Owing to the fact that an increased cellular force with the growing number of RBCs leads to larger traction, which has been reported by Isfahani and Freund (2012), CTC adhesion may be inhibited. In fact, a larger CTC velocity than the average blood velocity is expected to prevent the firm adhesion in CTC metastasis.



Fig. 8 Ratio of CTC velocity to the average blood velocity for different microvessels at various hematocrits

The effect of the CTC on the blood flow resistance was quantified by introducing the relative flow resistance, which is the ratio of the computed apparent viscosity with a CTC to the apparent viscosity without a CTC (Fedosov and Gompper 2014). Here, the computed apparent viscosity is defined by $\eta_{app} = \pi \Delta P D^4 / 128 Q L$. Figure 9 presents the relative flow resistance for blood flow in microvessels with a CTC. Generally, the presence of CTC increases the flow resistance, resulted from the increasing volume fraction of cells. This effect is more pronounced in smaller microvessels because of the larger relative volume fraction of CTC compared to that in large microvessel. In smaller microvessels, the larger probability of CTC adhesion induced by the stronger CTC confinement at high hematocrit leads to a larger flow resistance. However, in microvessels of 20µm, the flow resistance is larger at *Hct* =0.1 due to CTC has a larger adhesion probability (Fig. 7), which is consistent with Fig. 6.



Fig.9 Relative flow resistance for blood flow in different microvessels with a CTC at various hematocrits

3.5 Dependence on RBCs aggregation at different flow rates

It should be noted that the above findings are based on the condition of low flow rates and without considering the RBCs aggregation. Under such a low flow rate, the RBCs scatter in the flow without RBCs aggregation, which is not consistent with the real blood. In this section, the effects of flow rate combined with RBCs aggregation on the adhesion of the CTC in the microvessel ($D = 20\mu m$) were investigated.

Firstly, for the case without RBC aggregation, by comparing the variation of the number of receptorligand bonds over the time period of around 0.8s at different flow rates, Figure 10 shows that at high flow rate (γ =17.72s⁻¹), the initial adherent CTC detaches from the wall quickly due to the increasing lift force. But after a short time, the RBCs flowing towards the vessel center expel the CTC from the RBC core and further initiate the adhesion of the CTC to the vessel wall, as shown in Fig. 11c. In addition, the number of formed bonds is more than that at the low flow rate due to the large deformation resulting from the large shear deformation, as plotted in Fig. 10b. The CTC maintains its original spherical shape at the low flow rate as the value of $L_T/D_T = 1$



Fig. 10 Variation of number of receptor-ligand bonds (a) and the deformation of CTC along the flow direction (b) for the cases of RBC aggregation and non-aggregation at different flow rates

The main effect of RBCs aggregation is that the time of CTC contact to the vessel extends and the presence of RBCs aggregation provides an additional wall-directed force to compress the adherent CTC (Fig. 11b and Fig. 11d), leading to an increasing number of receptor-ligand bonds. At the low flow rate (γ =2.2s⁻¹), the RBC aggregation enables the CTC to stably roll along the surface of the vessel wall with a certain number of bonds. Nevertheless, the weak deformation inhibits the formation of firmed adhesion. However, the large deformation induced by the high flow rate increases the CTC contact area, thus the number of bonds increases considerably, which can be seen at *t* = 0.4s for the case of RBC aggregation in Fig. 10. If the adhesive force is larger than the hydrodynamic force exerted on the CTC, firm adhesion may be formed.

(a)

(b)



Fig. 11 Snapshots of the flow of a CTC and RBCs for the cases of (a) $\gamma = 2.2s^{-1}$, without RBC aggregation, (b) $\gamma = 2.2s^{-1}$, with RBC aggregation, (c) $\gamma = 17.72s^{-1}$, without RBC aggregation, (d) $\gamma = 17.72s^{-1}$, with RBC aggregation

4 Conclusions

The effects of flowing RBCs on the adhesion of CTC in microvessels were examined in terms of the hematocrit, vessel size, flow rate and RBC aggregation. At the low flow rate, increasing the hematocrit can detach the initial adherent CTC from the vessel wall due to the enhanced lift force in larger microvessels. While an inverse effect was found in smaller microvessels. This is because in a stronger confinement, a growing wall-directed force on the CTC is expected at higher hematocrit. Also, the velocity of CTC is larger than the average blood velocity in smaller microvessels but the ratio of CTC velocity to the mean blood velocity drops to around 1 in larger microvessel and even the CTC flows slower than the blood stream at *Hct* =0.1 as the CTC nearly moves near the vessel wall. In addition, the presence of CTC increases the blood resistance, which is more pronounced for the case of CTC adhesion. Lastly, the strong tendency of RBCs migrating towards the vessel in higher flow rate enables the detached CTC to contact with the vessel wall again. And the induced large deformation increases the number of receptor-ligand bonds for the adherent CTC. By adding the RBCs aggregation, an enhanced CTC adhesion can be found because it enables the CTC to stably roll along the vessel wall at the low flow rate. And an additional wall-directed force further compresses the CTC, leading to an increasing number of receptor-ligand bonds.

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