

DIVERGENCE-CONFORMING AND FULLY-IMPLICIT SIMULATION OF MICROSCLAE BLOOD FLOW

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ABSTRACT

In the last decades, modeling of cell-scale blood flow has attracted remarkable attention in computational mechanics. In vessels with a diameter smaller than 500 μm , blood cells are explicitly included in the model to study various phenomena in which the own motion of individual blood cells is of interest. This leads to a fluid-structure interaction (FSI) problem in which both blood cells and blood plasma are incompressible. We have recently extended our immersed method for FSI [1-3] in order to utilize divergence-conforming B-splines for the spatial discretization while maintaining a fully-implicit time discretization. Divergence-conforming B-splines have already been proven to increase the accuracy and robustness of immersed FSI methods in [4]. We have also developed a scalable implementation of our method which is necessary to tackle the application of cell-scale blood flow. Our fluid solver is based on the scalable block-preconditioning strategy developed in [5].

We will show the effects of size, deformability, and density when it comes to isolate nucleated cells using mechanical means. This could help to improve existent medical devices to isolate circulating tumor cells from blood samples.

Finally, we will study the flow patterns and rheology of blood in normal and stenosed arterioles. The stenosis of arterioles can lead to important diseases as coronary microvascular disease and lacunar infarct. The flow characteristics of stenosed arterioles are quite different from those of macrovascular stenosis due to the fact that blood behaves as a Newtonian fluid in large arteries, but as a non-Newtonian fluid in arterioles.

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