

Multiscale Thrombosis Simulations on Massively Parallel Computers

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We have developed a multiscale simulator for the primary stage of thrombus formation as a result of a platelet adhesion to an injured vessel wall. The simulator numerically treats fluid-membrane interactions at the continuum level [1][2], and also ligand-receptor interactions between proteins at the molecular level [3]. From the simulated results obtained using a massively parallel computer, we discuss the enhancement of the thrombus formation in view of the particulate flow dynamics.

REFERENCES

- [1] Ii, S., Gong, X., Sugiyama, K., Wu, J., Huang, H. and Takagi, S. (2012) A full Eulerian fluid-membrane coupling method with a smoothed volume-of-fluid approach, *Comm. Comput. Phys.*, **12**: 544–576.
- [2] Ii, S., Sugiyama, K., Takagi, S. and Matsumoto, Y. (2012) A computational blood flow analysis in a capillary vessel including multiple red blood cells and platelets, *J. Biomech. Sci. Engrg.*, **7**: 72–83.
- [3] Shiozaki, S., Ishikawa, K.L. and Takagi, S. (2012) Numerical study on platelet adhesion to vessel walls using the kinetic Monte Carlo method, *J. Biomech. Sci. Engrg.*, **7**: 275–283.