

Micelle Separation In Microfluidic Devices

Sunjin Song^{a)}, Min Sun Yeom^{*b)}, Kyung Hyun Ahn^{a)}, Seung Jong Lee^{a)}

a) School of Chemical and Biological Engineering, Seoul National University,
Shillim9-dong, Gwanak-gu, Seoul 151-744, Korea

b) Supercomputing Application Support Team,
Korea Institute of Science and Technology Information (KISTI),
373-1 Guseong-dong, Yuseong-gu, Daejeon, 305-701 Republic of Korea
E-mail:msyeom@kisti.re.kr

Key Words: Brownian Dynamics, Hydrodynamic Interaction, Microchannel

ABSTRACT

The lab-on-a-chip-based systems have significant possibility for many life science applications, including genomic analysis, and cell counting and characterization. Integrated microfluidic devices use diffusion-based separation and detection, which allows the parallel flow in a single microchannel. In order to predict the behavior of a blood cell in a long microchannel, Brownian dynamics simulations with the bead-spring model were carried out for a micelle. We model a micelle that is an aggregate of the amphiphilic molecules in order to consider a blood cell. Each amphiphilic molecule is represented through a chain of several effective monomers. All the effective monomers are connected through a finitely extensible nonlinear elastic (FENE) potential. The diffusion tensor in this work includes the hydrodynamic interaction tensor, which relates the velocity perturbation and is obtained by finite element interpolation.