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This paper was published online on 26 March 2007 with an incorrect version of Fig. 3. The figure has been corrected as of 15 May 2007. The figure is incorrect in the printed version of the journal; therefore, the figure has been published below for the benefit of the print readership.

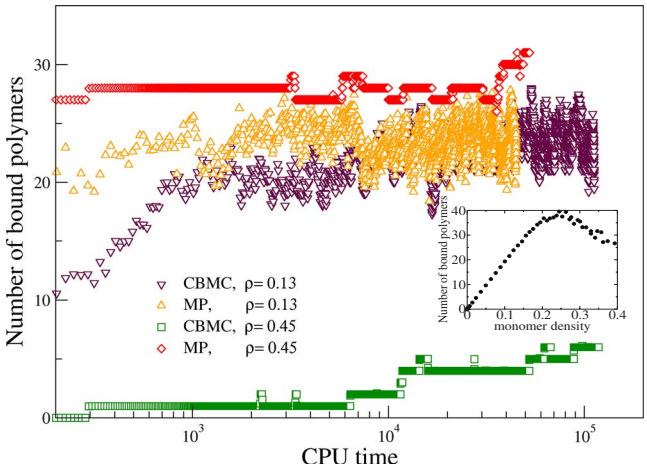

FIG. 3. (Color online) Density dependence of the equilibration rate of the MP and CBMC schemes. The figure shows the number of bound (bridge or loop) polymers between two walls as a function of CPU time. As in Fig. 2, the spacing between the two walls is 8 lattice units. The fraction of binding sites on the surface, $\rho_s=0.02$ and the binding energy, $\epsilon_b=10k_BT$. Results are shown for a high monomer concentration (45%) and a low monomer concentration (13%). The inset shows the density dependence of the number of bound polymers.

