

S2 Table: Variation of scaling exponent of MSD

Type	Theory	Simulation LJ strength ϵ	Fits to simulation ² $\phi = 0.4$	Fits to simulation ³ (unconfined)
Disconnected noninteracting beads	$\lambda = 1$	$\epsilon = 0$	$\lambda = 1$	
Disconnected interacting beads [1]	$\lambda = 3/4$	$\epsilon = 0.25, 0.5$	$\lambda = 0.8$	
Phantom polymer (Rouse dynamics [2])	$\lambda = 1/2$	$\epsilon = 0$	$\lambda = 0.5$	$\lambda = 0.5$
Polymer in good solvent ¹	$\lambda = 6/11$	$\epsilon = 0.25$	$\lambda = 0.4$	$\lambda = 0.5$
Polymer in poor solvent ¹	$\lambda = 2/5$	$\epsilon = 0.5$	$\lambda = 0.4$	$\lambda = 0.25$
Polymer melts (reptation [3])	$\lambda = 1/4$	$\epsilon = 1$		$\lambda = 0.25$

¹ Polymer of length M that is confined in a space has scaling exponent of MSD $\lambda = \frac{2\nu}{2\nu+1}$, where ν is related with MSD $\sim M^{2\nu}$ [4].

² S7 Fig

³ S5 Fig

Table: MSD, $\langle r^2 \rangle \sim \tau^\lambda$

References

- [1] Bénichou O, Illien P, Oshanin G, Sarracino A, Voituriez R. Diffusion and subdiffusion of interacting particles on comblike structures. Physical review letters. 2015;115(22):220601.
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- [4] Liu L, Shi G, Thirumalai D, Hyeon C. Chain organization of human interphase chromosome determines the spatiotemporal dynamics of chromatin loci. PLoS computational biology. 2018;14(12):e1006617.