Efficient simulation of Fractional Brownian Motion for several values of the Hurst exponent

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Fractional Brownian motion

- Efficient algorithm for long walks
- Distribution of endpoints

Fractional Brownian Motion

Translocation of polymer through pore: viral injection of DNA DNA sequencing with engineered channels s(t): position of chain at time t s(t > 0) = 0: absorbing boundary Proposal [Zoia, Rosso, Majumdar, PRL 2009]: Described by fractional Brownian motion = Gaussian process with $\langle s(t_1)s(t_2)\rangle \sim t_1^{2H} + t_2^{2H} - |t_1 - t_2|^{2H}$ $\Rightarrow C(t_1 - t_2) = \langle [s(t_1) - s(t_2)]^2 \rangle \sim |t_1 - t_2|^{2H}$ *H*: Hurst exponent: H = 1/2: Brownian motion, H > 1/2: correlation, H < 1/2: anticorrelation $(H = 1/(1 - \nu))$, where $R_a \sim N^{\nu}$ [Chuang, Kantor, Kadar 2001]) Rescaled variable: $v(t) = s(t)/t^{H}$ Prediction for $y \to 0$: $P(y) \sim y^{\phi}$ with $\phi = (1 - H)/H$

Random Walks

Traditional method for (non-absorbed) walks of length L

- 1. Vector ξ of $\tilde{L} \ge L$ Gaussian random numbers ξ_i
- 2. For (approximate) correlation: Fourier transform
- 3. Create walk $s(t) = \sum_{i < t} \xi_i$
- 4. Accept if $s(t) \ge 0$ for all t (non absorbed)
- Problem: Success probability of non absorbance (*persistence*) $\sim t^{-\phi}$





s(t) /

Monte Carlo approach

Basic idea:

Markov chain of vectors $\xi^{(0)} \rightarrow \xi^{(1)} \rightarrow \xi^{(2)} \rightarrow \dots$ step: change fraction of $\xi^{(l)}$ accept if walk not absorbed



Possible: additional reweighting $w \sim y^{\kappa}$ ($\kappa = -\phi \rightarrow$ "flat" sampling near y = 0)





Testcase pure Brownian motion (H = 0.5)

Distribution exactly known [Zoia, Rosso, Majumdar, PRL 2009]

 $P(y) = y \exp(-y^2/2)$



Superdiffusive case (H = 2/3)



 \rightarrow prediction $\phi = (1 - H)/H = 1/2$ well found $(y \rightarrow 0)$ confirmed medium-scale behavior y^{γ} ($\gamma > \phi$) predicted by [Wiese, Rosso, Majumdar, PRE 2012]

Subdiffusive case (H = 1/4)



 \rightarrow strong finite-length effects converges towards prediction $\phi = (1 - H)/H = 3 \ (y \rightarrow 0)$ in contrast to prediction $\phi = 2$ [Amitai, Kantor, Kadar, PRE 2010] (simulation of effective model for N = 257 coupled particles)



- Fractional Brownian motion: translocation of polymers
- Using large-deviation/MC approach: walks $L = 10^7$ feasible
- (Angti-)correlations readily included
- Reweighting: focus to region of interest \rightarrow better statistics
- $\phi(H = 0.25) \gg 2$: in favor of Zoia et al prediction

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