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Supplemental Information

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Behavior**

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Cytoplasmic RNA-Protein Particles Exhibit Non-Gaussian Subdiffusive Behavior: Supplemental Information

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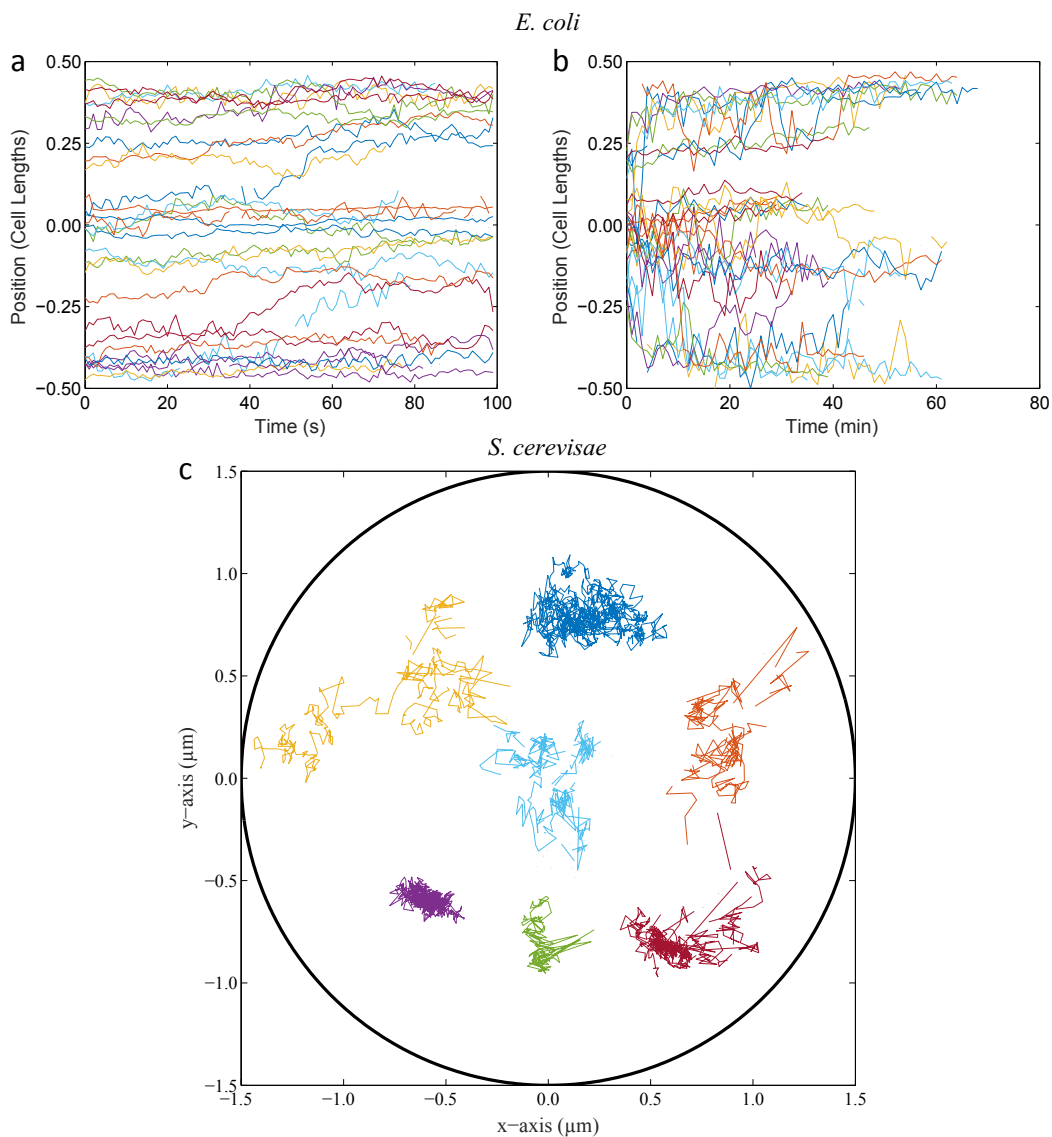


FIG. 1: **Representative trajectories of RNA-protein particles.** (ab) Sample trajectories of RNA-protein particles diffusing in the *E. coli* cytoplasm for (a) 1 s and (b) 1 min intervals between position measurements. (c) Sample trajectories placed arbitrarily in a circle with a diameter of 3 μm (typical size of an *S. cerevisiae* cell).

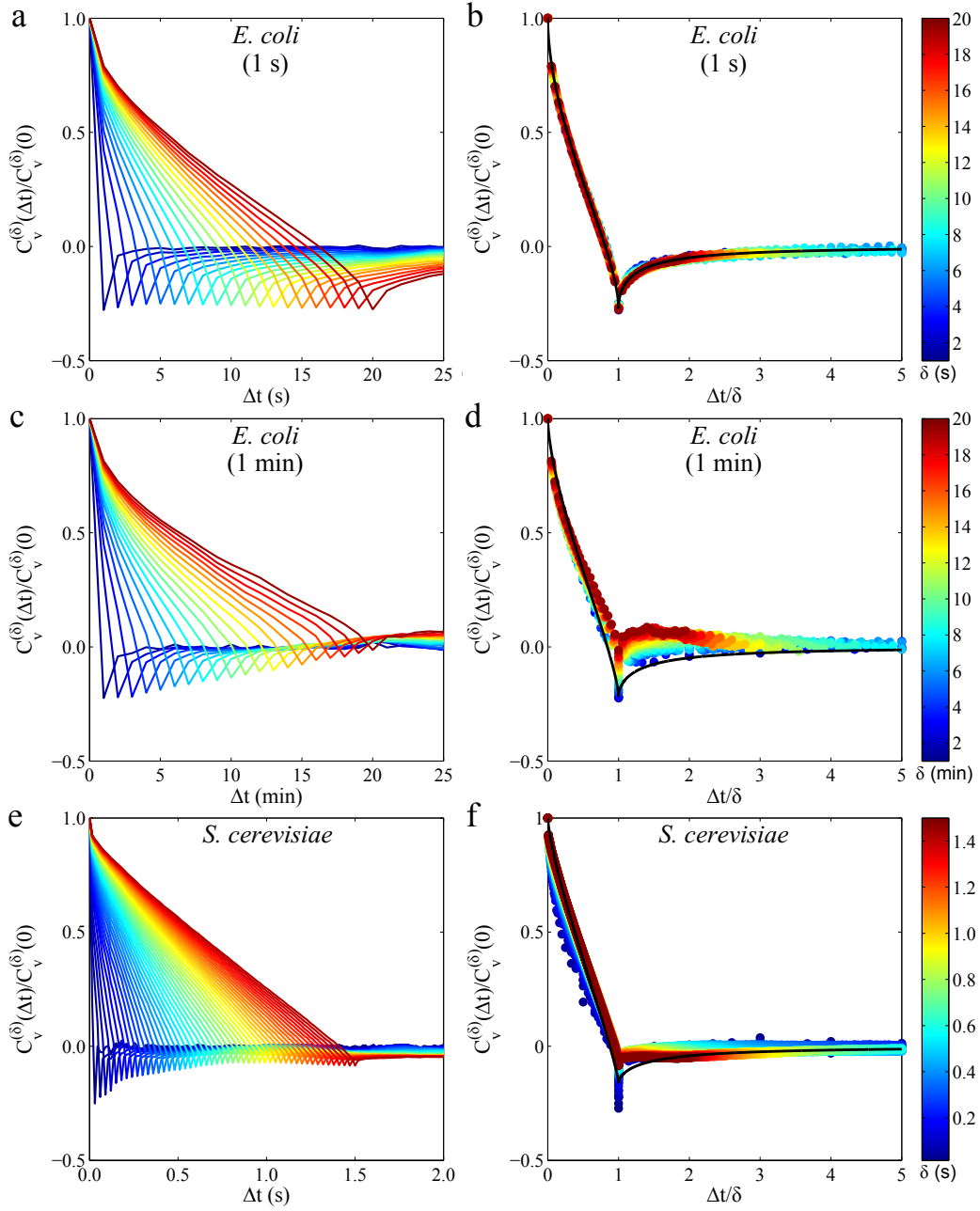


FIG. 2: **RNA-protein particles exhibit viscoelastic behavior.** (a-d) Velocity autocorrelation function of RNA-protein particles in *E. coli* for particle position measurements taken at one second intervals (a,b) and one minute intervals (c,d) [1] and corrected for drift due to cell growth. (e,f) Velocity autocorrelation function of RNA-protein particles in *S. cerevisiae* [2]. The velocity autocorrelation function is defined as $C_v^{(\delta)}(\Delta t) = \langle v_\delta(\Delta t) \cdot v_\delta(0) \rangle$ where the velocity $v_\delta(t) = [x(t + \delta) - x(t)]/\delta$ for the discrete time interval δ . (b), (d), and (f) are the same as (a), (c), and (e) respectively but with a differently scaled x-axis to show that the negative correlation peak occurs at $\Delta t/\delta = 1$. The black lines in (b), (d), and (f) are theoretical predictions for fractional Brownian motion with the measured values of α from the eMSD inserted into the equation $[(\eta+1)^\alpha + |\eta-1|^\alpha - 2\eta^\alpha]/2$, where $\eta = \Delta t/\delta$ [3].

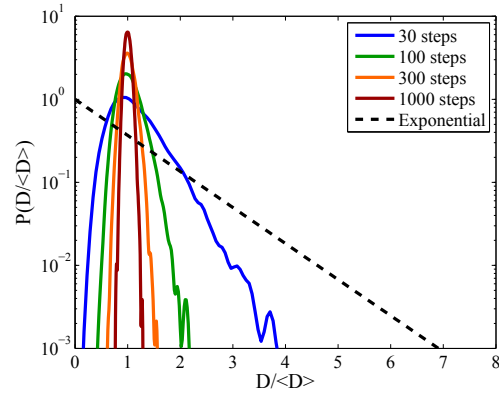


FIG. 3: **The effect of track length on the diffusivity distribution.** The distribution of diffusivities normalized by their mean for 10,000 fractional Brownian motion simulations of varying track lengths. Diffusivities are calculated using a power law fit to the first 15 points of each trajectory's time-averaged MSD. Distributions are plotted using a kernel density estimation with a Gaussian kernel and width determined by Silverman's rule of thumb.

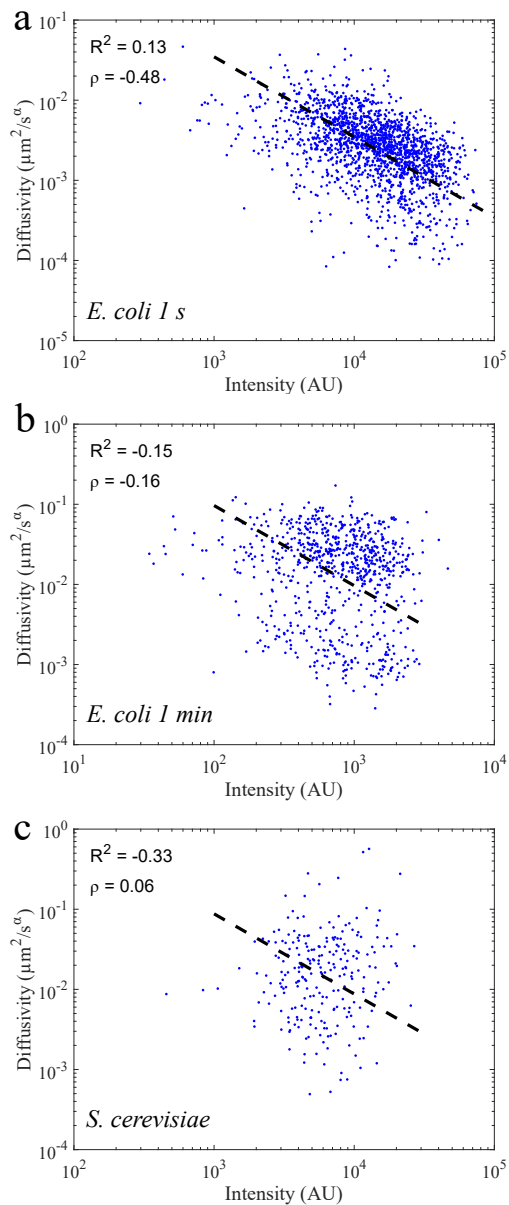


FIG. 4: **The relation between probe diffusivity and fluorescence intensity.** Particle diffusivities as a function of their mean intensity for the *e. coli* 1 s (a), *E. coli* 1 min (b), and the *S. cerevisiae* data (c). The black dashed line represents a best fit of the model $y = A/x$ with fit parameter A . We show R^2 and Pearson correlation coefficient (ρ) for each model fit to the data on a loglog scale.

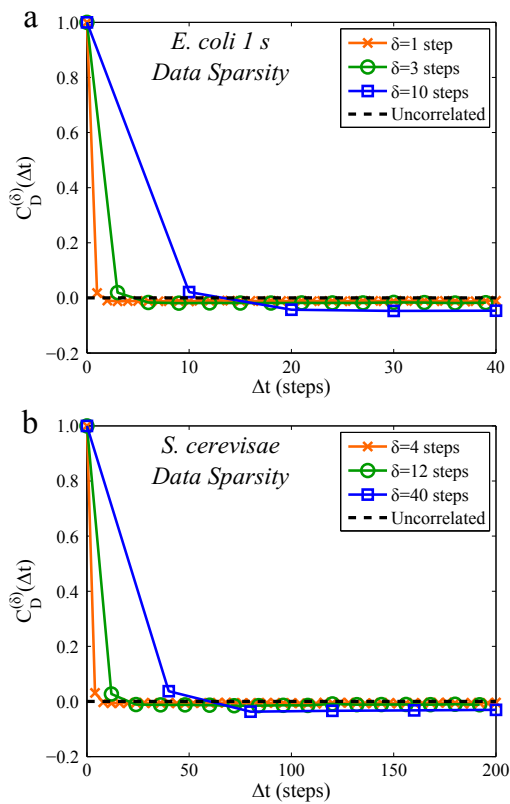


FIG. 5: **The effect of finite track lengths on the diffusivity autocorrelation function.** Diffusivity autocorrelation function for fractional Brownian motion simulations that have the same data sparsity and 10x the size of the *E. coli 1 s* data set (a) and the *S. cerevisiae* (b).