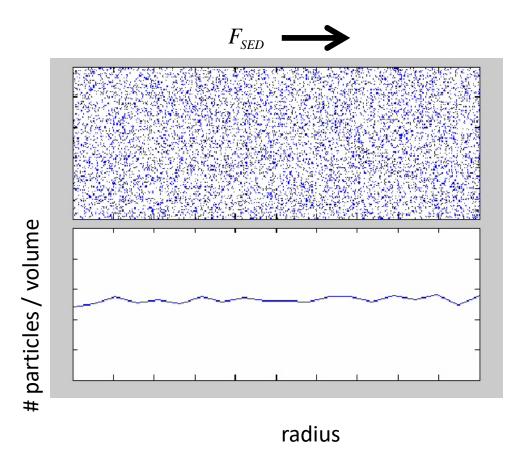
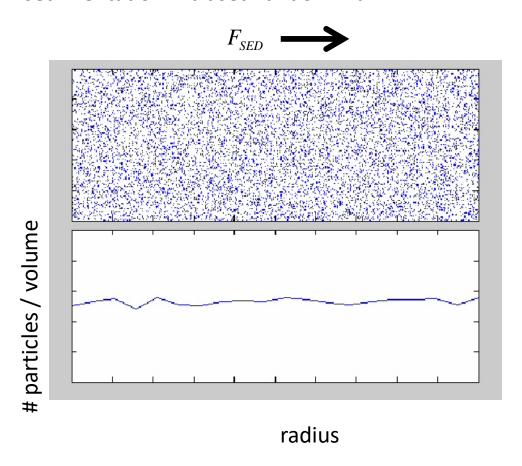
sedimentation = biased random walk



- → real-time detection systems triggered with revolution of rotor
- → allows to measure sedimentation, diffusion, polydispersity

$$M(1-\overline{v}\rho) = \frac{s}{D}RT$$

sedimentation = biased random walk



→ after long time, thermodynamic equilibrium reports on buoyant molar mass

$$c(r) = c(r_0) \exp^{\frac{M(1-\bar{v}\rho)}{RT}\frac{1}{2}(r^2 - r_0^2)}$$

for study of protein complexes

