

## Questions about theory and simulation

1. Why do you set a correlation time  $\tau$  before simulation, then use the simulation results to calculate  $\tau$  again? Do you expect them to be different?

au, the time scale of active noise, lies in the heart of the theory and simulation. The following velocity updating rule in your simulation,

$$u_{i_n} = u_{i_{n-1}} - \nu_i u_{i_{n-1}} + \nu_i \sqrt{B_i dt} N_{ni}, \quad i = x, y, z.$$

has to be designed in such a way, that velocity u is exponentially correlated,  $\langle u(t)u(t')\rangle \propto e^{-|t-t'|/\tau}$  (although I don't know how this updating rule enforces this correlation).

Simulation returns xyzt data, you use them to calculate the correlation time again:

saturation part of the MSD,ie,  $\lim_{t\to\infty} \langle \Delta x(t)^2 \rangle + \langle \Delta y(t)^2 \rangle$ . To compare we use  $\tau$  and  $r = r_o - r_i$  to make dimensionless the quantities from the experiments and also compute  $\tau$  for each simulation to make  $\hat{t}$  dimensionless. And

Would you expect them to be different? (The only thing that can make  $\tau$  different is the reflection boundary condition, and only very frequent occurance can make the difference significant enough to be noticed, IMO)

2. Theoretically, would you expect au to be a property of a bacterial suspension and independent of how you measure it?

We've been trying to measure  $\tau$  in different ways, e.g. with 2- $\mu$ m particles and with oil droplets as tracers. At higher concentration, we can also measure the flow persistence by PIV to get a  $\tau$ . Are these  $\tau$ 's expected to be the same (theoretically)?

Looking at the assumptions made in the simulation, any object immersed in an active bath exhibits a noisy motion with exponentially correlated velocity. Hence, I would infer that at least the tau measured from 2- $\mu$ m particles should apply to oil droplets. Do you think this is a way to test the assumption?

3. A similar question as Q#1, there are two quantities very similar to each other,  $\Pi_3$  and  $d_{st}$ . The former is an input to the simulation and the latter is the result. How are they different?

$$\Pi_3 = \Pi_1 \Pi_2 = \frac{v\tau}{r}, \quad \text{Bacteria versus confinement},$$
(1.23)

$$d_{st} = \frac{v_{tc}\tau}{r},\tag{1.24}$$

Again the only thing that can distinct the two is the boundary condition.