Bioinformatician with the
Monash Genomics and Bioinformatics Platform.

▶ R workshops for biologists with big datasets.

▶ Learning to process large data in blocks, with `arrow`, parquet.

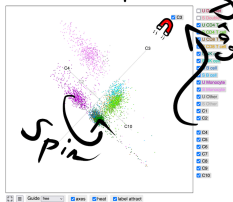▶ Wondering what is UMAP doing to my data.

scRNA-seq

Figure 1: I put an underdamped Langevin dynamics physics engine in a
Javascript tour widget (`langevitour`). What else could we put it into?
Diffusion-based generative models?