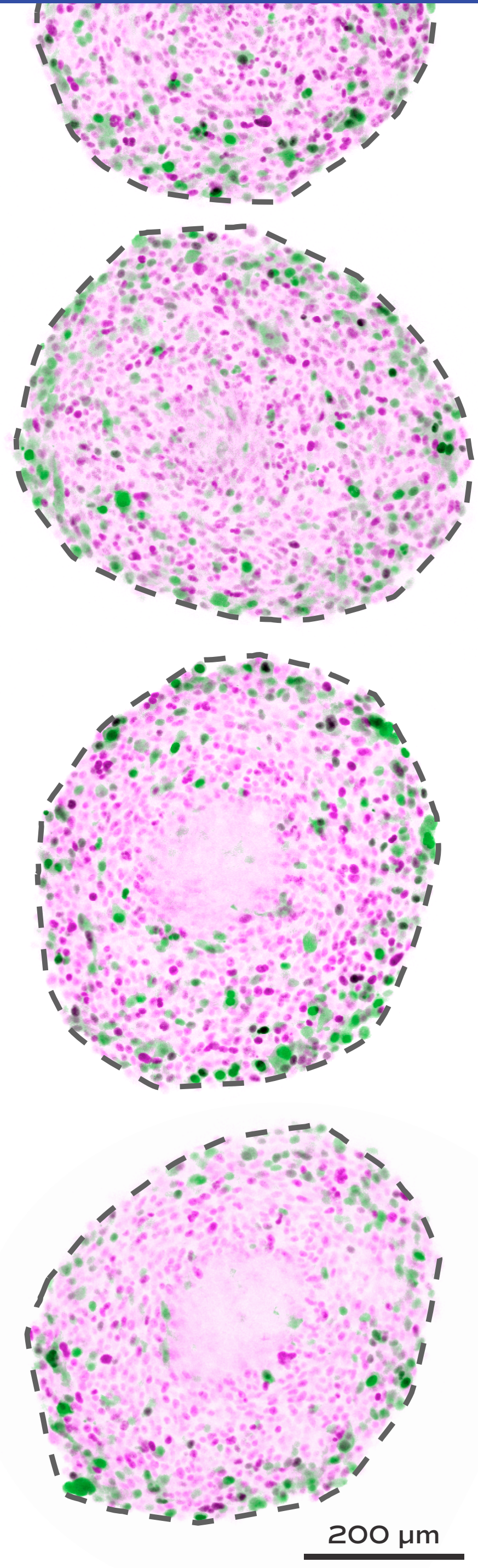
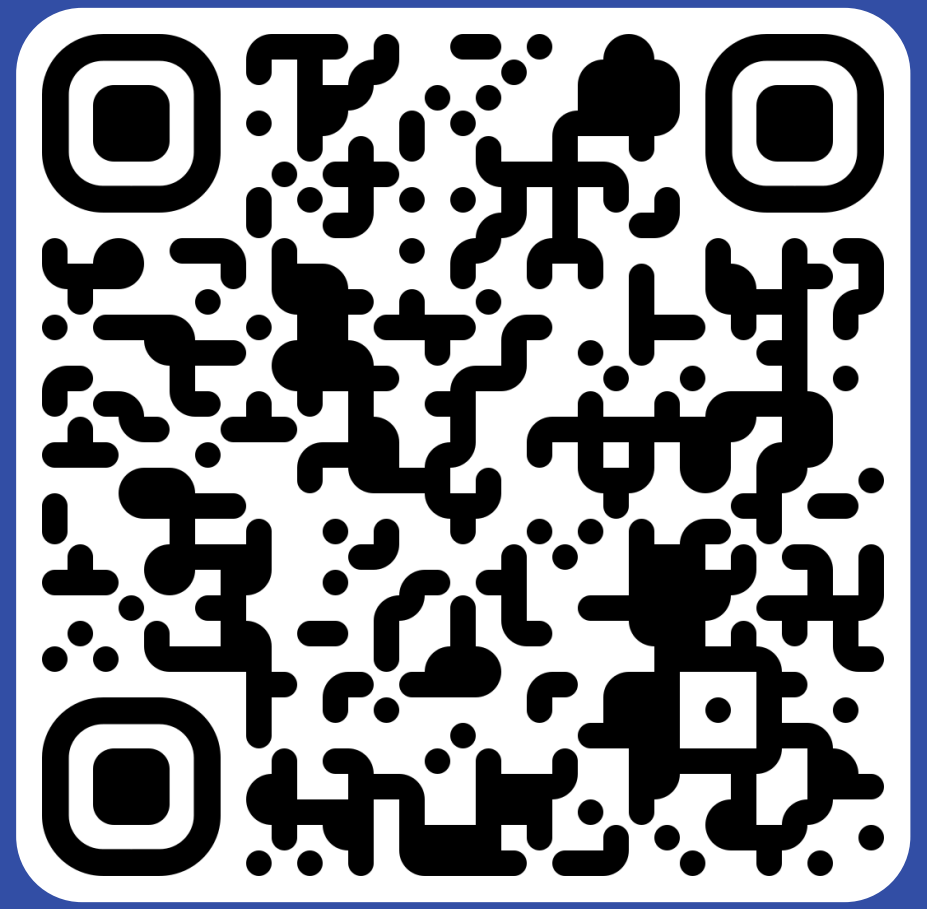


Modelling heterogeneity with random parameters

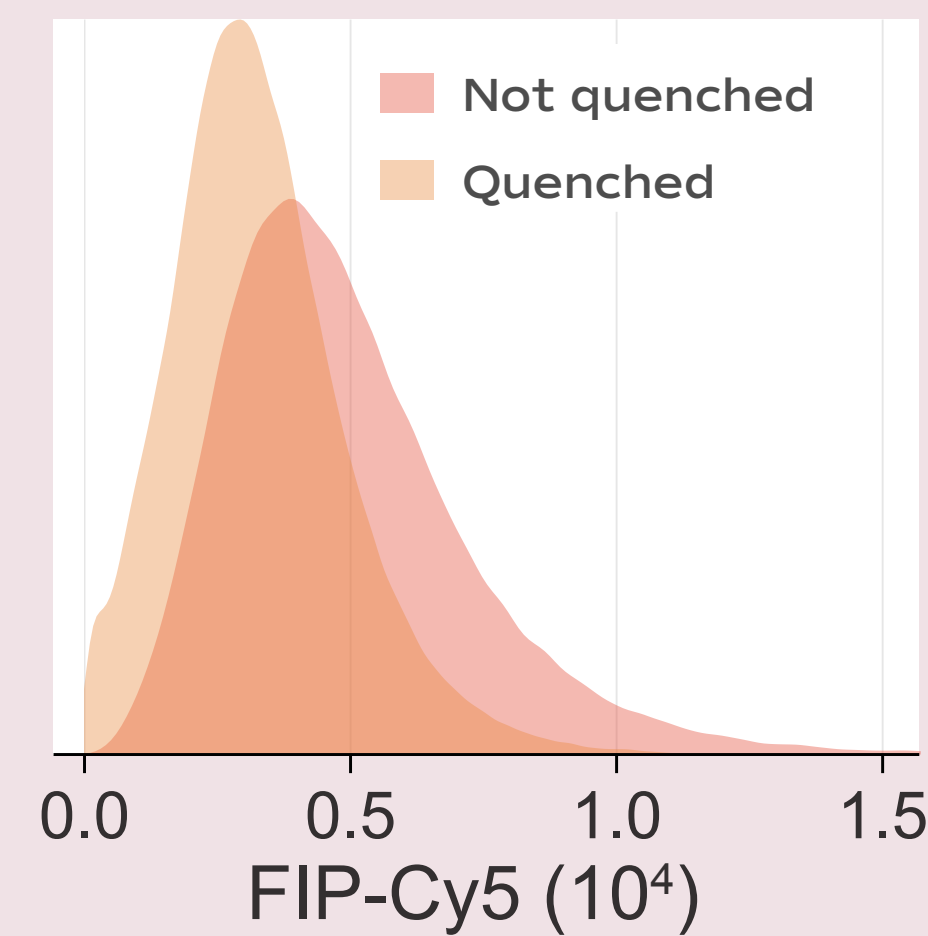
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Heterogeneity in biology

► **Figure 1**
Flourescent measurements relating to amount of internalised molecules in ~10 000 C1R cells.



◄ **Figure 2**
Identically prepared tumour spheroids from an *in vitro* experiment. Melanoma cells are induced with fluorescent cell cycle indicators (green indicates proliferation, magenta indicates cells in gap 1).

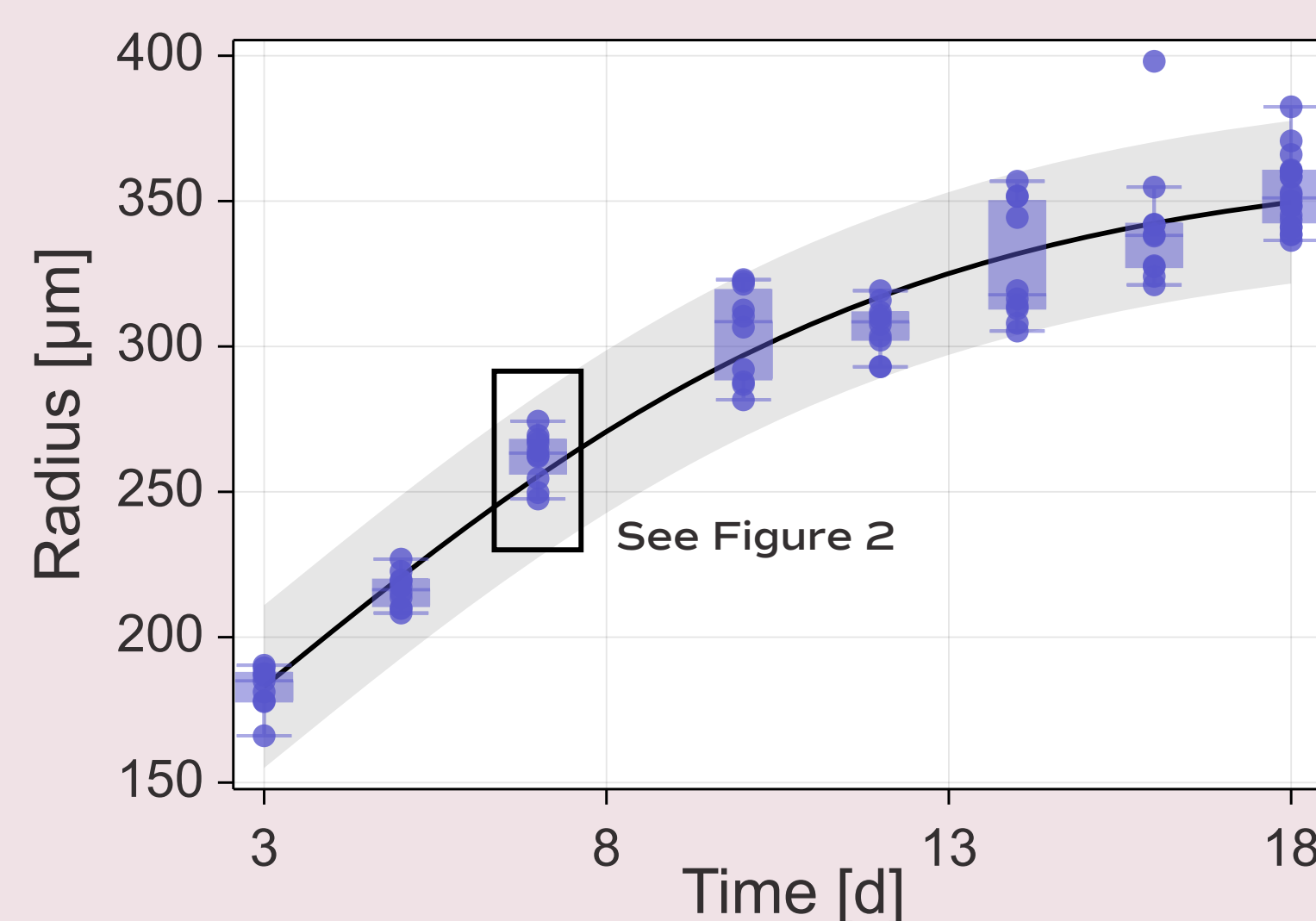
- **Biological variability** is seen even in **tightly controlled experiments** (figs. 1 and 2).
- Such variability is routinely captured in quantitative analysis by coupling a **deterministic mathematical model** to an independent **probabilistic observation process**.
- For example, consider coupling observations of tumour spheroid radius, $r^{\text{obs}}(t)$, to the logistic growth model using normal noise:

$$\frac{dr(t)}{dt} = \lambda r(t) \left(1 - \frac{r(t)}{R}\right), \quad r(0) = r_0.$$

$$r^{\text{obs}}(t) = r(t) + \varepsilon, \quad \varepsilon \sim \mathcal{N}(0, \sigma^2).$$

- The model is capable of recapturing experimental observations (fig. 3).

But what about variability in the initial condition and the dynamical parameters?



◄ **Figure 3**
Experimental measurements of temporal tumour spheroid radius described using the logistic growth model with additive normal noise. Shown is experimental data (blue) and model prediction (mean and 95% prediction interval).

Dealing with parameter variability

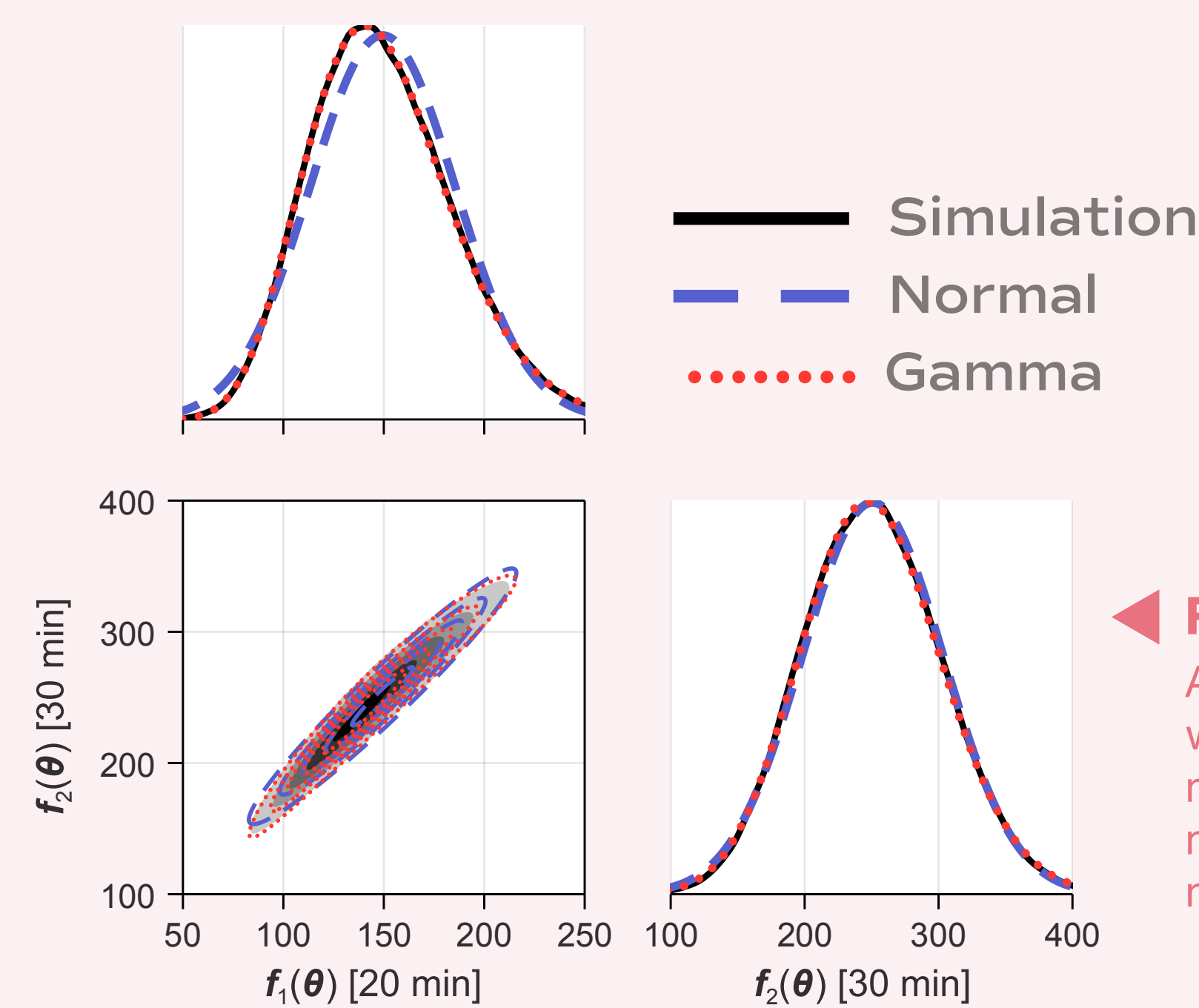
- We aim to better capture potential sources of variability by describing parameters and measurements as **random variables**.
- We take a fast, novel **direct approach** and construct an approximate output distribution for a given model and set of input (parameter) distributions.
- Consider that the mathematical model maps inputs (i.e., parameters), θ , to a set of outputs, $f(\theta)$. Then we wish to approximate the distribution of $f(\theta)$.

One-dimensional example

- Consider that both input and output are one-dimensional, then **expectation of the Taylor expansion** of $f(\theta)$ about $\hat{\theta} = \langle \theta \rangle$ yields

$$\langle f(\theta) \rangle \approx f(\hat{\theta}) + \underbrace{\frac{df(\hat{\theta})}{d\theta} \langle \theta - \hat{\theta} \rangle}_{=0} + \frac{1}{2} \underbrace{\frac{d^2f(\hat{\theta})}{d\theta^2} \langle \theta - \hat{\theta} \rangle^2}_{=\text{Var}(\theta)}.$$

- We can do this for inputs and outputs of **any dimension** yielding **analytical approximations for the mean, covariance, and skewnesses** of $f(\theta)$ in terms of the moments of θ .
- We then approximate the distribution of $f(\theta)$ using normal and gamma distributions with the same moments (fig. 4).



◄ **Figure 4**
Approximate solution to the logistic model with random parameters using a moment-matched normal distribution (blue; two moments) and gamma distribution (red; three moments).

Parameter inference

- We can **directly construct a likelihood** using our approximate output distribution, which describes the data distribution for a given set of input distributions and allows application of any likelihood-based inference tool (here, we use profile likelihood).

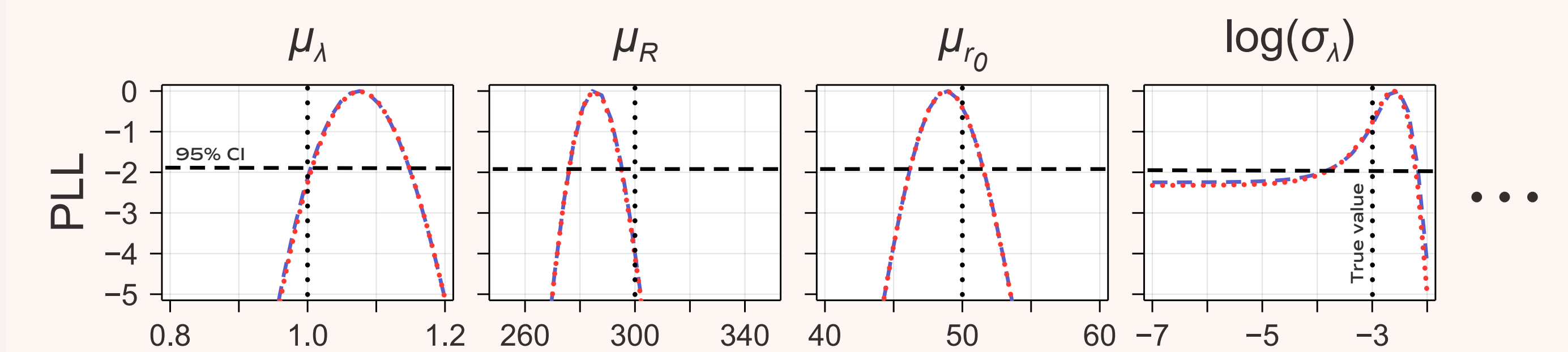
Random parameter logistic model

- Consider the logistic model with parameter distribution

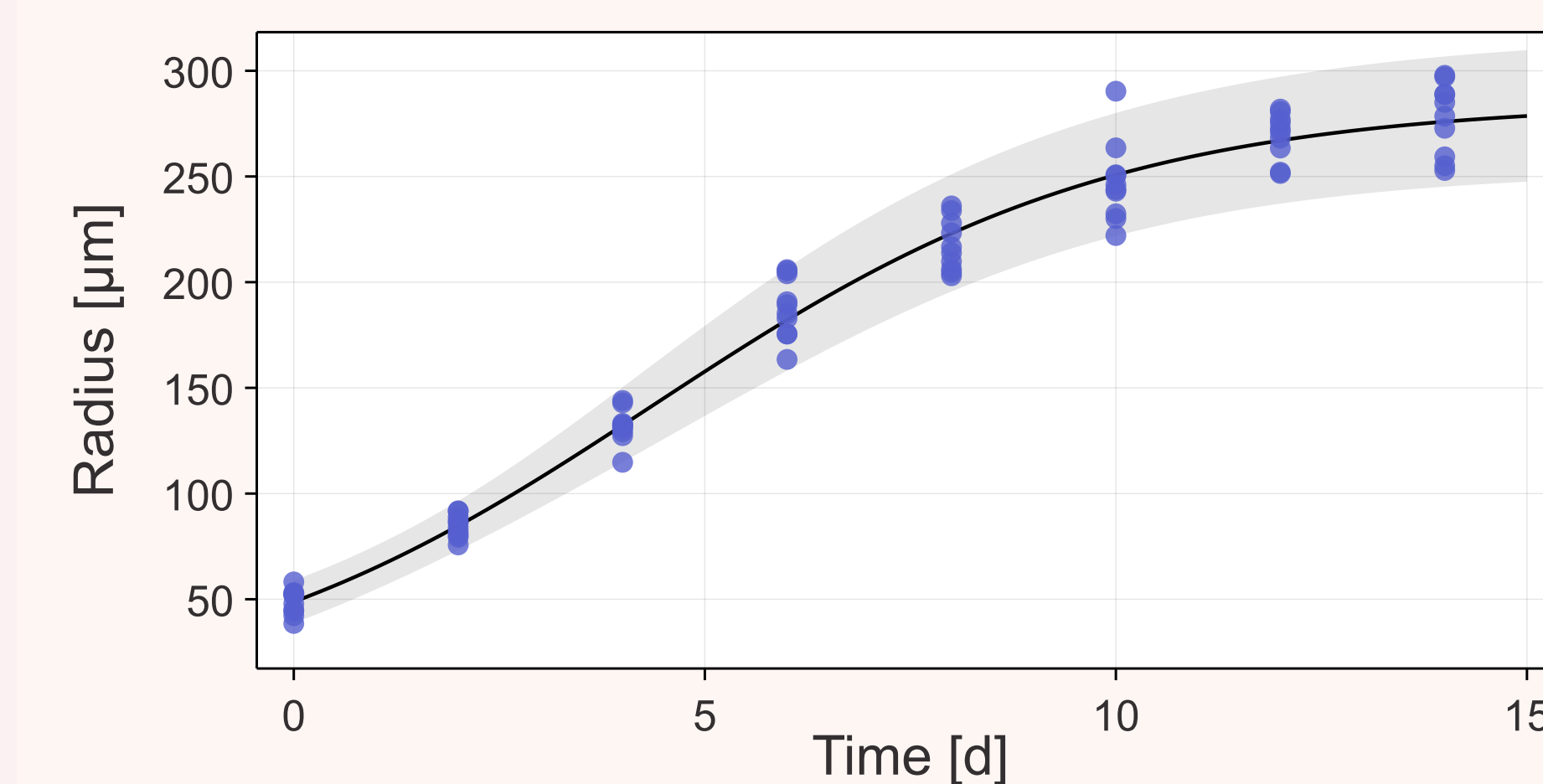
$$r_0 \sim \mathcal{N}(\mu_{r_0}, \sigma_{r_0}^2), \quad \lambda \sim \mathcal{N}(\mu_\lambda, \sigma_\lambda^2),$$

$$R \sim \mathcal{N}(\mu_R, \sigma_R^2), \quad \varepsilon \sim \mathcal{N}(0, \sigma_\varepsilon^2),$$

where $\xi = [\mu_{r_0}, \mu_\lambda, \mu_R, \ln \sigma_{r_0}, \ln \sigma_\lambda, \ln \sigma_R, \ln \sigma_\varepsilon]^\top$ is a vector of unknown “hyperparameters” that we wish to infer.



▲ **Figure 5**
Profile likelihoods for random parameter logistic model using synthetic data (fig. 6). Horizontal line shows the threshold for a 95% confidence interval.



◄ **Figure 6**
Synthetic data and predictive distribution (mean and 95% prediction interval) for the random parameter logistic model.

- In addition to inferring parameters and quantifying uncertainty, we much more accurately capture the variability in the data (fig. 6).
- **Julia module** for implementation can be applied to any differential-equation model, for a wide variety of input distributions (correlated, bimodal, skewed, etc).

