

Risk-averse design under uncertainty

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Risk-averse optimization of genetic circuits under uncertainty

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Problem statement

Proposed methodology

Example: design of a repressilator

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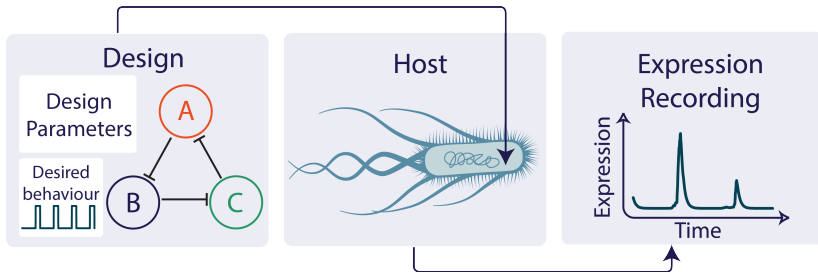
Example: design of a repressilator

Motivation

- ▶ Designing systems such that they have some prescribed properties is a ubiquitous problem.
- ▶ Mathematical/computer models of the system are often used to accelerate the design, e.g.
 - ▶ Hospitals: models are used to predict patient flow and optimise staffing.
 - ▶ Traffic management: models are used to optimise road networks and reduce congestion.
 - ▶ Synthetic biology: models are used to design genetic circuits.
- ▶ Design is framed as an optimisation problem.
- ▶ Issue: Models are not perfect and model-based designs may not achieve the predicted performance in reality.

Example: Design of a genetic circuit

- ▶ Genes form circuits that can be modelled as dynamical systems. Similar to circuits in electrical engineering.
- ▶ A network of three genes that inhibit (suppress) each other defines an oscillator, called "repressilator".
- ▶ Issue in genetic circuit design: performance on the computer may not be indicative of performance in-vivo.

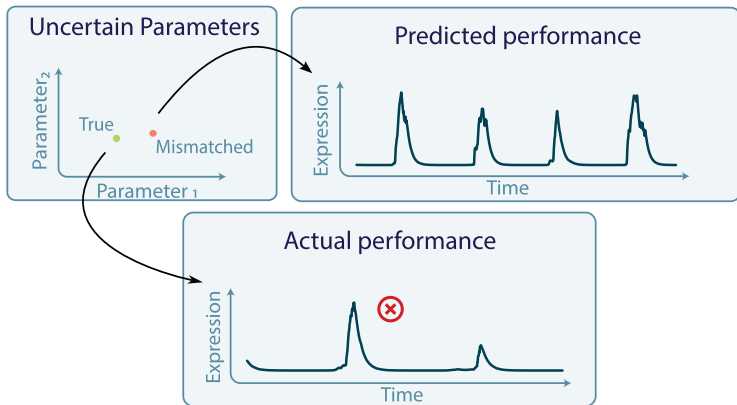


A case for design under uncertainty

- ▶ Hypothesis: gap between designed and actual performance is due to uncertainty (limited knowledge) about the real system.
- ▶ Two sources of uncertainty:
 1. Irreducible (aleatoric) uncertainty: modelled as system noise ω with distribution $p(\omega)$.
 2. Reducible (epistemic) uncertainty: modelled as system parameters η with prior distribution $p(\eta)$.
- ▶ Both types of uncertainty need to be considered in the design; otherwise we design the system under wrong assumptions.

A case for design under uncertainty

- ▶ Reconsider the example of the design of a repressilator.
- ▶ Assuming wrong values for the model parameters results in a design that works on the computer but not in reality.



Problem statement

- ▶ We assume we are given:

1. a stochastic generative model of the system:

$$\mathbf{x} = g(\omega, \eta, \phi), \quad \omega \sim p(\omega) \quad (1)$$

where \mathbf{x} describes the system, η are unknown system parameters not under our control, and ϕ are design parameters that we can control.

2. a prior belief $p(\eta)$
 3. data $\mathcal{D}_o = (\mathbf{x}_o, \phi_o)$ on the system from previous design attempts with ϕ_o
 4. a loss function $\tilde{L}(\mathbf{x}) = L(\omega, \eta, \phi)$ that measures how good \mathbf{x} is.
- ▶ Goal: Find designs η that are expected to work well even in unfavourable conditions.

Program

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Overview of our approach

Our approach consists of three steps:

1. Bayesian inference to update our belief about η given \mathcal{D}_o
2. Thompson sampling to obtain a set of promising designs.
3. Risk-management to select suitable designs.

Step 1: Bayesian inference

- ▶ Failed designs are oftentimes written off.
- ▶ We instead learn from them using Bayesian inference, updating prior to posterior belief $p(\boldsymbol{\eta}|\mathcal{D}_o)$.
- ▶ Depending on the model, we can compute the posterior with MCMC, variational inference, or simulation-based (likelihood-free) inference.
- ▶ Won't go further into this part of the pipeline.

Step 2: Thompson sampling

- ▶ The loss $\tilde{L}(\mathbf{x}) = L(\omega, \eta, \phi)$ measures how good a specific \mathbf{x} is.
- ▶ It is random due to the randomness of ω and η .
- ▶ Let us average over the system noise ω but pushforward the randomness of η .
- ▶ This results in a stochastic process with realisations

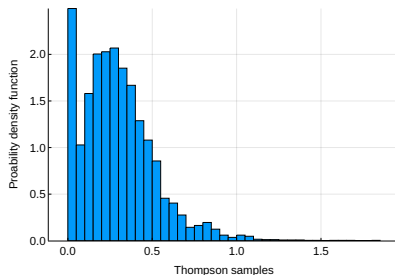
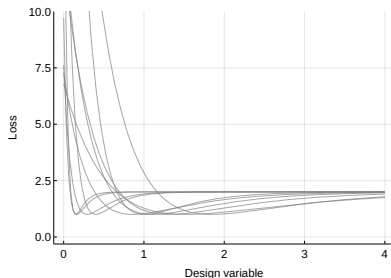
$$L_i(\phi) = \mathbb{E}_{\omega} [L(\omega, \eta_i, \phi)], \quad \eta_i \sim p(\eta | \mathcal{D}_o) \quad (2)$$

- ▶ Thompson samples ϕ_i are defined as $\phi_i = \operatorname{argmin}_{\phi} L_i(\phi)$
- ▶ They are random variables: Uncertainty about the system translates into uncertainty about the design.
- ▶ Named after William R Thompson (1887-1972).

(On the Likelihood That One Unknown Probability Exceeds Another in View of the Evidence of Two Samples. Biometrika (1933))

Simple example of Thompson sampling

- ▶ Let $x = g(\omega, \boldsymbol{\eta}, \phi) = \eta_1 \exp(-\phi \eta_2) + \omega$, with $\boldsymbol{\eta} = (\eta_1, \eta_2)$, $\phi \geq 0$, and the loss is the squared distance between x and 1.0.
- ▶ Gamma distributions for η_1 and η_2 , standard normal for ω .
- ▶ Left: Loss as a function of ϕ for different realisations of η_1 and η_2 .
- ▶ Right: Distribution of the Thompson samples



Step 3: Risk management

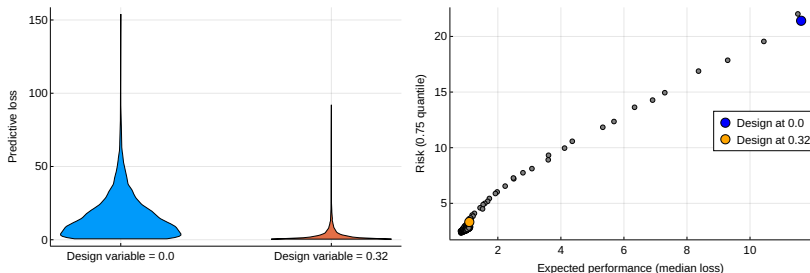
- ▶ Each Thompson sample ϕ_i has an associated predictive distribution of the loss,

$$L(\omega, \eta, \phi_i), \quad \omega \sim p(\omega), \eta \sim p(\eta|\mathcal{D}_o) \quad (3)$$

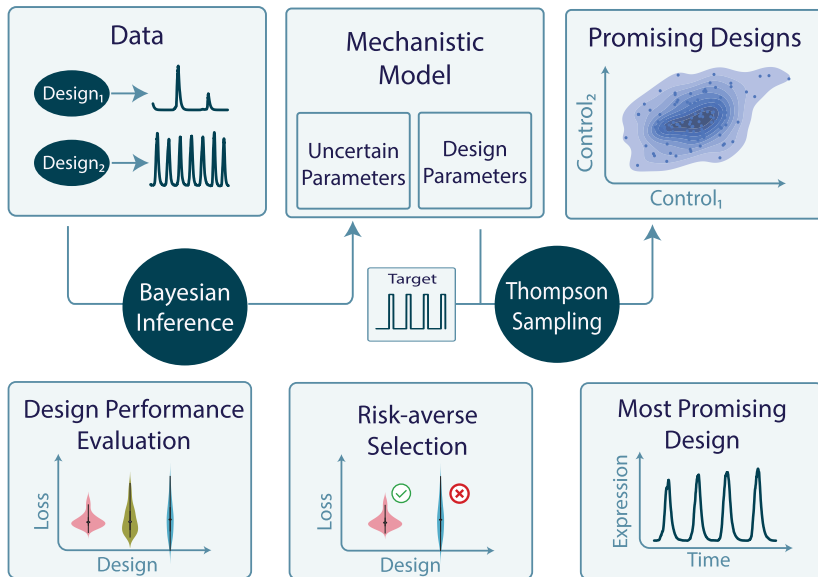
- ▶ Distribution reflects system noise and our uncertainty about the system.
- ▶ The distribution can be used to rank the Thompson samples.
- ▶ We reduce the distribution to two statistics:
 - ▶ Median as a robust, risk-neutral, measure of expected performance
 - ▶ 0.75 quantile as a measure of risk, measuring how much worse the performance is when performing worse than the median.
- ▶ Pick design that achieves desired trade-off.
- ▶ Other choices possible, e.g. for optimistic design.

Back to the example

- ▶ Left: Predictive loss for two Thompson samples
- ▶ Right: Mapping out expected performance vs risk for different values of ϕ .
- ▶ No trade-off between performance and risk in this example.



Summary of our approach



Program

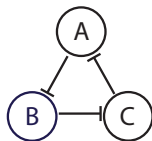
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Model

- ▶ The model $\mathbf{x} = g(\boldsymbol{\omega}, \boldsymbol{\eta}, \boldsymbol{\phi})$ is implicitly defined via the solution to a stochastic differential equation for the protein expressions p_1, p_2, p_3 .

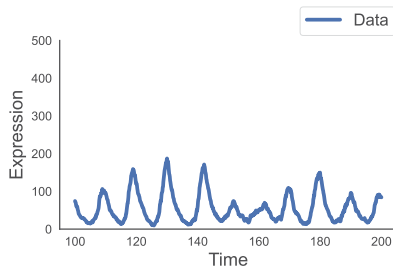
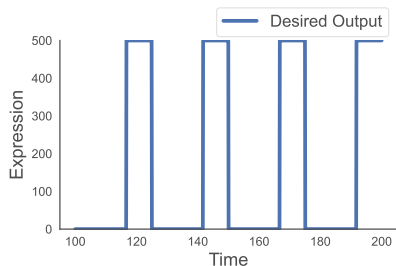


$$dp_i = \left(\underbrace{\frac{\alpha}{1 + \left(\frac{p_{i-1}}{K}\right)^n}}_{\text{Production}} - \underbrace{\gamma \cdot p_i}_{\text{Degradation}} \right) dt + \underbrace{\left(\sqrt{\frac{\alpha}{1 + \left(\frac{p_{i-1}}{K}\right)^n}} dW_{i1} + \sqrt{\gamma \cdot p_i} dW_{i2} \right)}_{\text{Noise}}$$

- ▶ Uncertain parameters $\boldsymbol{\eta} = (\gamma, n)$: degradation rate, Hill coefficient
- ▶ Design (controllable) parameters $\boldsymbol{\phi} = (\alpha, K)$: production rate and dissociation constant

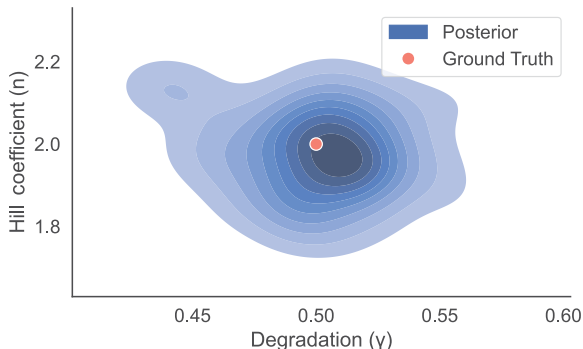
Design goal and data

- ▶ Design goal: 4 oscillations during 100 minutes (left).
- ▶ Failed attempt: $\phi = (334, 10)$. Used as data \mathcal{D}_o to update our belief about η (right)



Step 1: Bayesian inference

- ▶ Prior on η : Uniform on $(0, 10)$.
- ▶ Posterior obtained with SMC-ABC.
- ▶ For invariance to temporal shifts of the trajectories, we matched the absolute value of the Fourier spectrum of simulated and observed data.

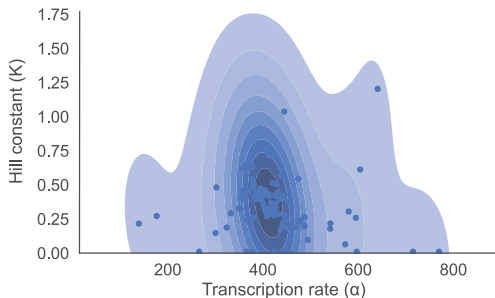


Step 2: Thompson sampling

- For each Thompson sample ϕ_i , we minimise the L_2 distance between the spectrum of the simulated and target trajectory.

$$L_i(\phi) = \mathbb{E}_{\omega} \left[\frac{1}{N} \sum_{n=0}^N (|F_{(\eta_i, \phi)}[n]| - |F_{target}[n]|)^2 \right], \quad \eta_i \sim p(\eta | \mathcal{D}_o)$$

- Distribution of Thompson samples.

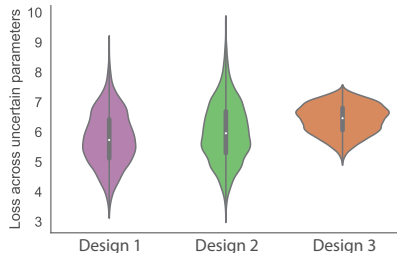
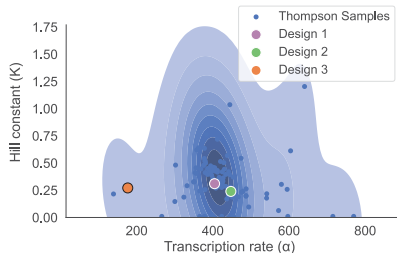


Step 3: Risk management

- ▶ To assess the Thompson samples ϕ_i , we evaluate their predictive loss distribution, using

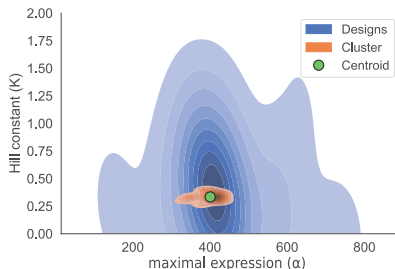
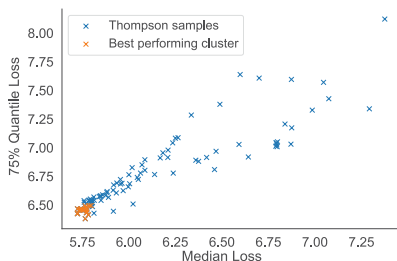
$$L(\omega, \eta, \phi_i), \quad \omega \sim p(\omega), \eta \sim p(\eta|\mathcal{D}_o) \quad (4)$$

- ▶ Examples



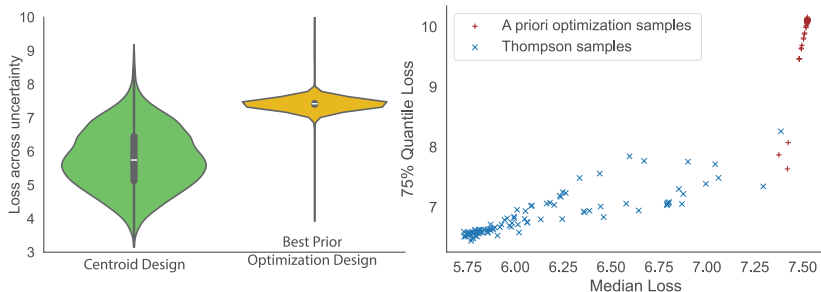
Step 3: Risk management

- ▶ We summarise each predictive loss distribution in terms of their median (expected performance) and 75% quantile (risk).
- ▶ Enables informed decision about suitable designs.
- ▶ Cluster marked in orange are most promising designs. Centroid as single best one.



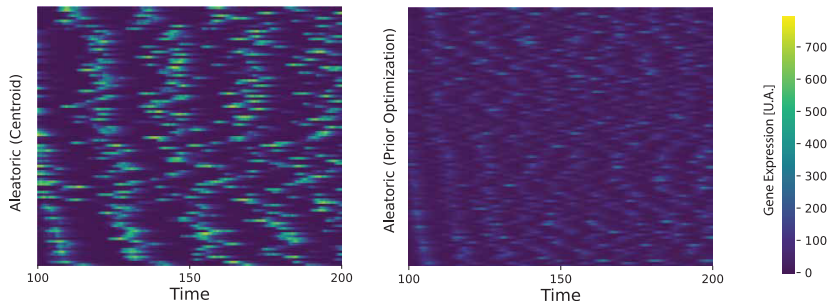
Evaluation

- ▶ Is there a benefit in using the failed design attempts to learn more about the systems?
- ▶ Comparison between Thompson samples for $\eta \sim p(\eta|\mathcal{D}_o)$ and $\eta \sim p(\eta)$ indicates large predicted performance difference.



Evaluation

- ▶ The centroid design point is *predicted* to be best. How good is it when deployed?
- ▶ Left fig: Design achieves four oscillations in the specified time-interval under ground truth η parameters.
- ▶ Right fig: This is not the case for best design under the prior.



Conclusions

- ▶ Talk was on designing systems with the help of models.
- ▶ Models are not perfect and model-based designs may not achieve the predicted performance in reality.
- ▶ Considered the case where the gap between designed and actual performance is due to uncertainty (limited knowledge) about the real system.
- ▶ Proposed an uncertainty-aware design methodology by combining:
 - ▶ Bayesian inference
 - ▶ Thompson sampling
 - ▶ Risk management
- ▶ Application in synthetic biology.
- ▶ Preprint with more results:
<https://www.biorxiv.org/node/4220646>

