# Efficient Approximation of the Likelihood for Complex Models

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## **Notation**

- ▶ Model a (random) function  $f: \Theta \to \mathcal{Y}$ .
  - ightharpoonup: parameter space
  - $ightharpoonup \mathcal{Y}$  : model output space
  - $ightharpoonup \mathbf{Y}_{\theta} := f(\theta)$  (assumed same form as  $\mathbf{Y}_{\text{obs}}$ ).





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- ► Y<sub>obs</sub> : a vector of observed data (incidence, prevalence, hospitalisations etc.)
- $S(\mathbf{Y}_{obs})$ : summary statistic (vector) of observed data (average weekly incidence etc.)





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  - ightharpoonup Or  $\mathcal{L}(\theta|S(\mathbf{Y}_{ ext{obs}})) := \Pr(S(\mathbf{Y}_{ ext{obs}})|\theta)$
- $\hat{m{ heta}} = \operatorname{arg\,max}_{m{ heta}} \mathcal{L}(m{ heta}|S(\mathbf{Y}_{\mathsf{obs}}))$





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- $\hat{\boldsymbol{\theta}} = \operatorname{arg\,max}_{\boldsymbol{\theta}} \mathcal{L}(\boldsymbol{\theta}|S(\mathbf{Y}_{\mathsf{obs}}))$
- $\blacktriangleright \ \mathsf{Pr}(\theta|S(\mathbf{Y}_{\mathsf{obs}})) \propto \mathsf{Pr}(S(\mathbf{Y}_{\mathsf{obs}})|\theta) \, \mathsf{Pr}(\theta)$





## Reality

- ► Explicit likelihoods often don't exist/are intractible
  - eg. agent based models





## A Standard Bayesian Solution

- Approximate Bayesian Computation (ABC)
  - 1. Sample  $\theta_i$  from prior
  - 2. Run model and observe  $\mathbf{Y}_{\theta_i}$
  - 3. Accept or reject  $\theta_i$  run based on how well  $\mathbf{Y}_{\theta_i}$  'matches'  $\mathbf{Y}_{\text{obs}}$ .





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- Discrepency function  $D: \mathcal{S} \times \mathcal{S} \to \mathbb{R}$ 
  - e.g. *p*-norm

$$||S(\mathbf{Y}_{\theta_i}) - S(\mathbf{Y}_{\mathsf{obs}})||_{p} := (\sum_{i=1}^{d} |S(\mathbf{Y}_{\theta_i}) - S(\mathbf{Y}_{\mathsf{obs}})|^{p})^{1/p}$$





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- $\triangleright \ \mathcal{D}(\theta) := D(S(\mathbf{Y}_{\theta}), S(\mathbf{Y}_{\text{obs}}))$





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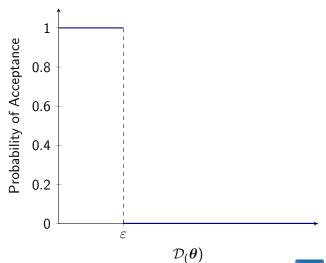
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- ▶ Rescale  $S(\cdot)$  appropriately (ie via a covariance matrix).
- $\triangleright \mathcal{D}(\theta_i)$  is 'how close' we were using parameters  $\theta_i$ .





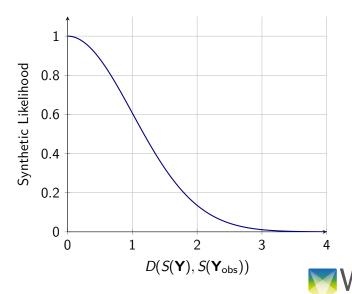
# Uniform Acceptance Probability







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## Overall Idea of my Research

▶ Can we predict  $\mathcal{D}(\theta_i)$  without having to evaluate  $f(\theta_i)$ ?





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- ▶ Can we predict  $\mathcal{D}(\theta_i)$  without having to evaluate  $f(\theta_i)$ ?
- Locally, hopefully yes.





## Gaussian Processes

- Random functions.
- Common examples Brownian motion, Ornstein Uhlenbeck process.





## Gaussian Processes on $\mathbb{R}^d$

## Definition (Gaussian Process)

A collection of random variables  $\{f(\mathbf{x})\}_{\mathbf{x}\in\mathbb{R}^d}$  is a Gaussian process if all finite dimensional distributions are multivariate normal distributed. That is, there is a function  $m: \mathbf{x} \to \mathbb{R}$  and kernel  $k: \mathcal{Y} \times \mathcal{Y} \to \mathbb{R}$  such that for all finite sets  $\{\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_n\}$ ,

$$\begin{bmatrix} f(\mathbf{x}_1) \\ f(\mathbf{x}_2) \\ \vdots \\ f(\mathbf{x}_n) \end{bmatrix} \sim \mathsf{MVN} \left( \begin{bmatrix} m(\mathbf{x}_1) \\ m(\mathbf{x}_2) \\ \vdots \\ m(\mathbf{x}_n) \end{bmatrix}, \mathbf{K} \right)$$

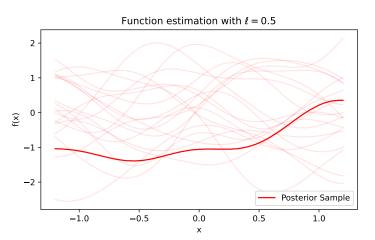
where

$$\mathbf{K} = \begin{bmatrix} k(\mathbf{x}_1, \mathbf{x}_1) & k(\mathbf{x}_1, \mathbf{x}_2) & \dots & k(\mathbf{x}_1, \mathbf{x}_n) \\ k(\mathbf{x}_2, \mathbf{x}_1) & \ddots & & \vdots \\ \vdots & & \ddots & \vdots \\ k(\mathbf{x}_n, \mathbf{x}_1) & \dots & \dots & k(\mathbf{x}_n, \mathbf{x}_n) \end{bmatrix}$$





# Gaussian Process Example Realisations







#### Covariance Kernel Motivation

- Kernel determines the amount of covariance between sets of indices.
- ▶ When the distance between indices is small, covariance needs to be large





## Common Covariance Kernels

Matern Kernel

$$k_{\nu}(x,x') = \sigma_k^2 \frac{2^{1-\nu}}{\Gamma(\nu)} \left( \frac{\sqrt{2\nu}||x-x'||}{\ell} \right)^{\nu} K_{\nu} \left( -\frac{\sqrt{2\nu}||x-x'||}{\ell} \right)$$

where  $K_{\nu}$  is a modified Bessel function (||  $\cdot$  || is the euclidean distance)

- $ightharpoonup \lfloor 
  u 
  floor$  times mean square differentiable.
- $u \to \infty$  infinitely mean square differentiable squared exponential covariance kernel (strong assumption)

$$k(x, x') = \sigma_k^2 \exp\left(-\frac{||x - x'||^2}{\ell}\right)$$





## Kernel Classes

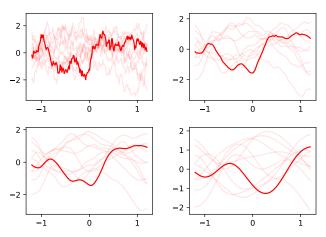


Figure: Matérn 1/2, 3/2, 5/2, and squared exponential kernels.





## Gaussian Process Regression

$$\begin{bmatrix} f(\mathbf{x}) \\ f(\mathbf{x}_*) \end{bmatrix} \sim \mathcal{N} \left( \begin{bmatrix} m(\mathbf{x}) \\ m(\mathbf{x}_*) \end{bmatrix}, \begin{bmatrix} \mathcal{K} & \mathcal{K}_* \\ \mathcal{K}_*^T & \mathcal{K}_{**} \end{bmatrix} \right)$$

implies

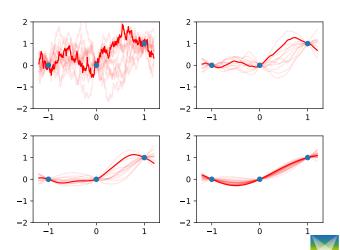
$$f(\mathbf{x})|f(\mathbf{x}_*) \sim \mathcal{N}\left(m(\mathbf{x}) + K_*K_{**}^{-1}(f(\mathbf{x}_*) - m(\mathbf{x}_*)), \ K - K_*K_{**}^{-1}K_*^T\right).$$





# Fitting our GP to data

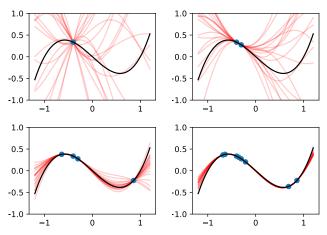
## GPs are 'priors'







# GP regression on x(x-1)(x+1)







## What if we have noise?

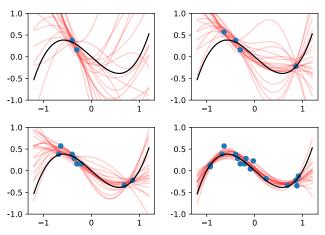
Add observation variance  $\sigma_o^2$ ,

$$\begin{bmatrix} f(\mathbf{x}_1) \\ f(\mathbf{x}_2) \\ \vdots \\ f(\mathbf{x}_n) \end{bmatrix} \sim \mathsf{MVN} \begin{pmatrix} \begin{bmatrix} m(\mathbf{x}_1) \\ m(\mathbf{x}_2) \\ \vdots \\ m(\mathbf{x}_n) \end{bmatrix}, \ \mathbf{K} + \sigma_o^2 \mathbf{I}_n \end{pmatrix}$$





# GP regression on $x(x-1)(x+1) + \epsilon$ , $\epsilon \sim (N(0, \sigma_o^2))$







## Overall Idea again

- lacktriangle Can we predict  $\mathcal{D}(m{ heta}_i)$  without having to evaluate  $f(m{ heta}_i)$  <2->
- ▶  $\mathcal{D}(\theta) \approx \mathcal{D}(\theta')$  for  $\theta$ ,  $\theta'$  close. <3->
- lacktriangle Approximate  $\mathcal{D}(m{ heta})$  by a Gaussian process  $\mathcal{D}_{\mathcal{GP}}(m{ heta})$





- ▶ High expected  $\mathcal{D}_{\mathcal{GP}}(\theta)$  with low variance = waste of time (and resources)
- Parameter Quantify this using a Bayesian acquisition function A, and choose  $\arg\min_{\theta} A(\theta)$





▶ Gutmann and Cor 2016 uses lower confidence bound

$$A_{\mathsf{LCB}}(\boldsymbol{\theta}) := \mu(\boldsymbol{\theta}) - \eta_t \sqrt{\mathbf{v}(\boldsymbol{\theta})}$$

- $ightharpoonup \mu(m{ heta}),\, {
  m v}(m{ heta})$  are posterior mean and variance
- ► (Claim of theoretical guarantees = load of rubbish)





Expected information

$$\begin{split} A_{\mathsf{EI}}(\boldsymbol{\theta}) := & \mathbb{E}(\min[\mathcal{D}_{\mathcal{GP}}(\boldsymbol{\theta}) - \mu_{\mathsf{min}}, 0)] \\ = & (\mu_{\mathsf{min}} - \mu(\boldsymbol{\theta})) \Phi\left(\frac{\mu_{\mathsf{min}} - \mu(\boldsymbol{\theta})}{\sqrt{\mathrm{v}(\boldsymbol{\theta})}}\right) \\ & + \sqrt{\mathrm{v}(\boldsymbol{\theta})} \phi\left(\frac{\mu_{\mathsf{min}} - \mu(\boldsymbol{\theta})}{\sqrt{\mathrm{v}(\boldsymbol{\theta})}}\right) \end{split}$$

- $ightharpoonup \mu_{\min} := \min_{\theta} \mu(\theta)$
- $lackbox{}{\hspace{-0.1cm}$\Phi$},\phi$  CDF and PDF of standard normal



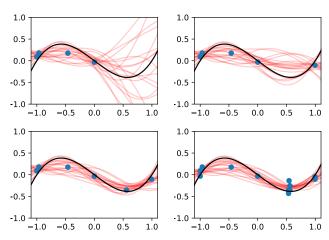


▶ Theoretical guarantees highly sensitive to choice of kernel





## Lower Confidence Bound

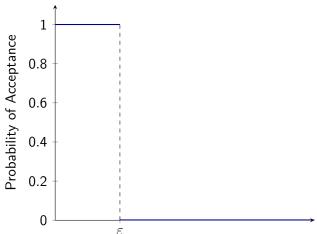






## Synthetic Likelihood

▶  $L(\theta|\mathbf{Y}_{\mathsf{obs}}) \approx P(\mathcal{D}_{\mathcal{GP}}(\theta) < \varepsilon)$  (up to a proportion)









#### Vivax Malaria

► Has dormant liver stage on top of blood stage infection that can cause relapse.





# Vivax Model - Champagne et. al

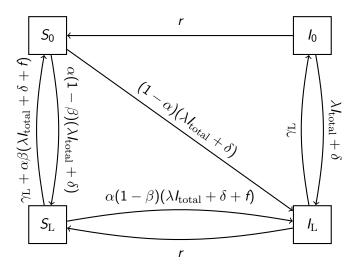




Figure: P. vivax model described by Champagne et al. 2022 WEH

# Ordinary Differential Equations - Champagne et. al

$$\begin{split} \frac{\mathrm{d}I_{\mathrm{L}}}{\mathrm{d}t} = & (1-\alpha)(\lambda I_{\mathrm{total}} + \delta)(S_0 + S_{\mathrm{L}}) + (\lambda I_{\mathrm{total}} + \delta)I_0 \\ & + (1-\alpha)fS_{\mathrm{L}} - \gamma_{\mathrm{L}}I_{\mathrm{L}} - rI_{\mathrm{L}} \\ \frac{\mathrm{d}I_0}{\mathrm{d}t} = & -(\lambda I_{\mathrm{total}} + \delta)I_0 + \gamma_{\mathrm{L}}I_{\mathrm{L}} - rI_0 \\ \frac{\mathrm{d}S_{\mathrm{L}}}{\mathrm{d}t} = & -(1-\alpha(1-\beta))(\lambda I_{\mathrm{total}} + \delta + f)S_{\mathrm{L}} + \alpha(1-\beta)(\lambda I_{\mathrm{total}} \\ & + \delta)S_0 - \gamma_{\mathrm{L}}S_{\mathrm{L}} + rI_{\mathrm{L}} \\ \frac{\mathrm{d}S_0}{\mathrm{d}t} = & -(1-\alpha\beta)(\lambda I_{\mathrm{total}} + \delta)S_0 + (\lambda I_{\mathrm{total}} + \delta)\alpha\beta S_{\mathrm{L}} + \alpha\beta fS_{\mathrm{L}} \\ & + \gamma_{\mathrm{L}}S_{\mathrm{L}} + rI_0 \end{split}$$





## Champagne Model Parameters

- ightharpoonup lpha : proportion of those infected but cleared of blood stage infections (through treatment)
- $\beta$ : a further proportion that are also cleared of liver stage parasites, given that they were also cleared of blood stage infection (radical cure)
- $\triangleright$   $\lambda$  : the rate of infection
- $ightharpoonup \gamma_L$  : rate of clearance of liver stage disease
- f: rate of relapse
- r: rate of blood stage clearance
- $ightharpoonup \delta = 0$  importation rate (fixed)





#### Model Calibration Data

▶ Doob-Gillespie algorithm with paper parameters reported in the original paper for 'observed data', 10 initial infections, 1000 people.



Expected information acquisition function



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    m obs}$ : weekly incidence around (stochastic) equilibrium
  - $ightharpoonup p_{\text{obs}}$ : prevalence around (stochastic) equilibrium
  - $ightharpoonup m_{\rm obs}$ : incidence in the first month of the epidemic



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$$\mathcal{D}(\alpha, \beta, \gamma_L, \lambda, f, r) = \ln \sqrt{\left(\frac{p - p_{\text{obs}}}{p_{\text{obs}}}\right)^2 + \left(\frac{m - m_{\text{obs}}}{m_{\text{obs}}}\right)^2 + \left(\frac{w - w_{\text{obs}}}{w_{\text{obs}}}\right)^2}$$

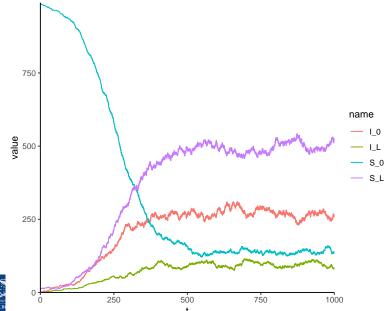
ightharpoonup (Log of the  $L_2$  norm of the relative differences)



Expected information acquisition function



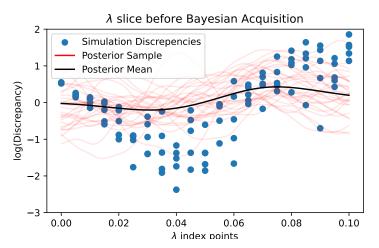
**Example Simulation** 





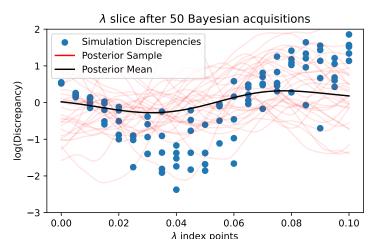






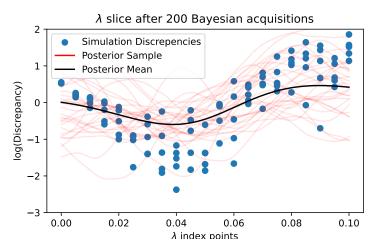






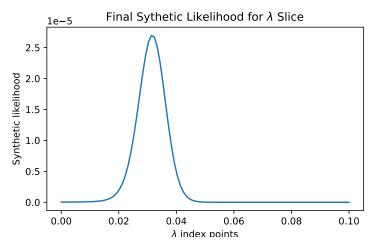






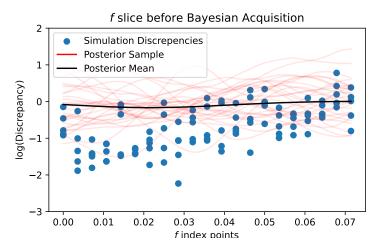






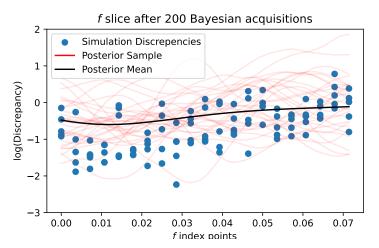






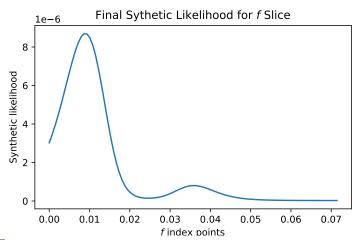
















#### Discussion

- Observation variance is considered constant across the GP (or log GP)
  - Particularly a problem at the threshold
- Assumes that normal/log-normal distribution approximates  $\mathcal{D}(oldsymbol{ heta})$
- Jumps where there is threshold/bifurcation behaviour
  - ► Student *t*—Process?





#### Thanks to

- ► Eamon Conway
- ▶ Jennifer Flegg



