Efficient Likelihood Approximation via Gaussian Processes

With an Application to a P. Vivax Malaria Model

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Introduction and Motivation

- ▶ 600,000 deaths/year, 75% children under 5
- Two main species *P. vivax* and *P. falciparum*
- P. falciparum main cause of death, but P. vivax historically underestimated.
- ▶ Proportion of *P. vivax* cases increased over last 50 years.





P. vivax has Dormant Stage

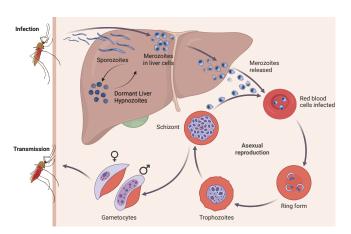


Figure: P. vivax lifecycle. Created with BioRender.com





Vivax Model - Champagne et. al

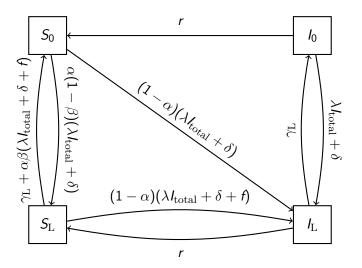




Figure: Champagne et al. 2022 P. vivax model



Champagne Model Parameters

- ightharpoonup lpha : proportion of those infected who clear blood stage infections through treatment
- β : proportion of those cleared of blood stage infection who are also cleared of liver stage parasites (radical cure)
- \triangleright λ : rate of infection
- $ightharpoonup \gamma_{\it L}$: rate of liver stage disease clearance
- f: rate of relapse
- r: rate of blood stage clearance
- $ightharpoonup \delta = 0$ importation rate (fixed)





The Problem

► How to calibrate model parameters?





The Problem

- ► How to calibrate model parameters?
- Simulations take long time (and models get a lot more complicated)





Notation

- ▶ θ vector of parameters e.g. $[\alpha, \beta, \gamma_L, \lambda, f, r]^T$
- ► Y_{obs}: a (summary) vector of observed data e.g. (weekly) incidence, prevalence, (monthly) hospitalisations





Notation

- θ vector of parameters e.g. $[\alpha, \beta, \gamma_L, \lambda, f, r]^T$
- ► Y_{obs}: a (summary) vector of observed data e.g. (weekly) incidence, prevalence, (monthly) hospitalisations
- **Y**_{θ}: a random vector of model statistics for given θ .





In an ideal world...

There would be an explicit form for the likelihood:

$$\mathcal{L}(oldsymbol{ heta}) := \mathsf{Pr}(\mathbf{Y}_{oldsymbol{ heta}} = \mathbf{Y}_{\mathsf{obs}} | oldsymbol{ heta})$$

- $\hat{\boldsymbol{\theta}} = \operatorname{arg\,max}_{\boldsymbol{\theta}} \mathcal{L}(\boldsymbol{\theta})$
- lacksquare $\mathsf{Pr}(m{ heta}|\mathbf{Y}_{\mathsf{obs}}) \propto \mathcal{L}(m{ heta})\,\mathsf{Pr}(m{ heta})$
- Off to the pub



Or not...

- Explicit likelihoods often don't exist/are intractible
 - Champagne model
 - Agent based models.





A Standard Bayesian Solution

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





What is 'matches'?

1.
$$\mathbf{Y}_{\theta_i} = \mathbf{Y}_{\text{obs}}$$





What is 'matches'?

Y_{θi} = Y_{obs}
 Good luck...





What is 'matches'?

- 1. $\mathbf{Y}_{\theta_i} = \mathbf{Y}_{\text{obs}}$
- 2. Rescale **Y**s, and use discrepency function $D: \mathbb{R}^d \times \mathbb{R}^d \to \mathbb{R}$ e.g. *p*-norm

$$D(\mathbf{Y}_{oldsymbol{ heta}_i}, \mathbf{Y}_{\mathsf{obs}}) := \left(\sum_{j=1}^d \left| \{\mathbf{Y}_{oldsymbol{ heta}_i} \}_j - \{\mathbf{Y}_{\mathsf{obs}} \}_j
ight|^p
ight)^{1/p}$$





Discrepency Function

 $\mathcal{D}(\theta) := D(\mathbf{Y}_{\theta}, \mathbf{Y}_{\text{obs}})$ how 'close' our model is to the observed data using parameters θ





ABC

- 1. Sample θ_i from prior
- 2. Run model
- 3. Accept θ_i if $\mathcal{D}(\theta_i) < \varepsilon$.





Overall Idea of my Research

- ▶ ABC fixes one problem but leaves another:
 - ▶ Don't need $\mathcal{L}(\theta)$.
 - ightharpoonup Evaluating $\mathcal{D}(heta)$ takes as long as a model run.





Overall Idea of my Research

- ► ABC fixes one problem but leaves another:
 - ▶ Don't need $\mathcal{L}(\theta)$.
 - lacktriangle Evaluating $\mathcal{D}(m{ heta})$ takes as long as a model run.
- $\triangleright \mathcal{D}(\theta), \mathcal{D}(\theta')$ will be highly correlated when θ is near θ' .
 - Gaussian Processes





Gaussian Process Setup

Formally we can assume that

$$Cov(\mathcal{D}(\theta_i), \mathcal{D}(\theta_j)) = k(\theta_i, \theta_j)$$

for some covariance kernel k that decays to 0 as θ_i is further away than θ_j .

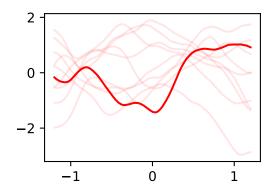




Gaussian Processes on \mathbb{R}^d

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.







Gaussian Process Continuity

▶ Induce continuity by forcing $k(\mathbf{x}, \mathbf{x}') \to \operatorname{Var}(f(\mathbf{x}))$ (hence $\operatorname{Cor}(f(\mathbf{x}), f(\mathbf{x}')) \to 1$) as $\mathbf{x} \to \mathbf{x}'$.





Common Covariance Kernels

- Choice of kernel determines smoothness
- ▶ Matérn Kernel with hyperparameter ν : $\lfloor \nu \rfloor$ times mean square differentiable.
- $u \to \infty$: infinitely mean square differentiable squared exponential covariance kernel (strong assumption)

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





k Determines Class of Functions

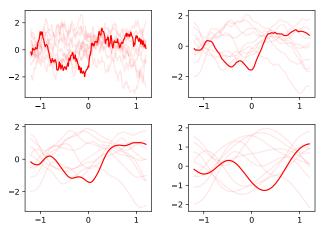


Figure: Matérn 1/2, 3/2, 5/2, and squared exponential kernels.





Kernel Hyperparameters

- Matérn and squared exponential kernel can both be written in the form $k(\mathbf{x}, \mathbf{x}') = \sigma_k^2 \kappa(||\mathbf{x}, \mathbf{x}'||/\ell)$
- ▶ $1/\ell$ rate of covariance decay



Discrepency Function Context

▶ Long term play: replace $\mathcal{D}(\theta)$ with a Gaussian process surrogate model approximation.





Discrepency Function Context

- ▶ Long term play: replace $\mathcal{D}(\theta)$ with a Gaussian process surrogate model approximation.
- ▶ What if we have observations already?





Gaussian Process Regression

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

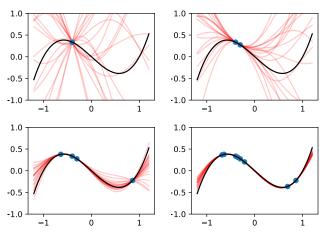
implies

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





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







Normal observation noise

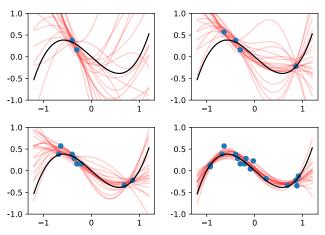
If observations actually $f(\mathbf{x}_i) + \varepsilon_i$, with $\varepsilon_i \sim N(0, \sigma_o^2)$ i.i.d., then

$$\begin{bmatrix} f(\mathbf{x}_1) + \varepsilon_1 \\ \vdots \\ f(\mathbf{x}_n) + \varepsilon_n \end{bmatrix} \sim \mathsf{MVN} \left(\begin{bmatrix} m(\mathbf{x}_1) \\ \vdots \\ m(\mathbf{x}_n) \end{bmatrix}, \, \mathbf{K} + \sigma_o^2 \mathbf{I}_n \right)$$





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







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Key Idea

- ▶ If $\mathbb{E}[\mathcal{D}(\theta)]$ can be well approximated by a Gaussian process and ...
- $ightharpoonup \mathcal{D}(\theta)$ approximately distributed $\mathit{N}(\mathbb{E}[\mathcal{D}(\theta)], \sigma_o^2)$ then...
- Approximate $\mathcal{D}(\theta)$ with $\mathcal{D}_{\mathcal{GP}}(\theta)$, a Gaussian process with observation noise.





ABC

- 1. Sample θ_i from prior
- 2. Run model
- 3. Accept θ_i if $\mathcal{D}(\theta_i) < \varepsilon$.





Approximate ABC...??

- 1. Sample θ_i from prior
- 2. Run model Simulate $\mathcal{D}_{\mathcal{GP}}(\theta_i) \stackrel{d}{\approx} \mathcal{D}(\theta_i)$
- 3. Accept θ_i if $\mathcal{D}_{\mathcal{GP}}(\theta_i) < \varepsilon$.





Synthetic Likelihood

Pr drawing and accepting heta using 'approximate' ABC is

$$\Pr(\mathcal{D}_{\mathcal{GP}}(\boldsymbol{\theta}) < \varepsilon) \Pr(\boldsymbol{\theta})$$

and hence

$$\hat{L}(oldsymbol{ heta}) := \mathsf{Pr}(\mathcal{D}_{\mathcal{GP}}(oldsymbol{ heta}) < arepsilon) pprox c\mathcal{L}(oldsymbol{ heta})$$

for some *c*.





Log Gaussian Process

▶ Alternatively we can model $\ln \mathcal{D}(\theta)$ as a Gaussian process $d_{\mathcal{GP}}(\theta)$.





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Log Gaussian Process

- Alternatively we can model $\ln \mathcal{D}(\theta)$ as a Gaussian process $d_{\mathcal{GP}}(\theta)$.
- Key assumptions becomes:
 - ▶ $\mathcal{D}(\theta)$ approximately distributed $LN(\cdot, \sigma_o^2)$.

$$\hat{L}(\boldsymbol{\theta}) := \Pr(d_{\mathcal{GP}}(\boldsymbol{\theta}_i) < \ln \varepsilon) \approx \Pr(\mathcal{D}(\boldsymbol{\theta}_i) < \varepsilon)$$





Where to sample $\mathcal{D}(\boldsymbol{\theta})$

- ▶ To generate a reliable $\mathcal{D}_{\mathcal{GP}}$, we need to sample widely
- ▶ Generating $\mathcal{D}(\theta)$ still costly...
- Therefore sample where:





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- Therefore sample where:
 - $ightharpoonup \mathbb{E}[\mathcal{D}(oldsymbol{ heta})]$ small,
 - $\triangleright \mathcal{D}(\theta)$ highly unknown.





Bayesian Acquisition Functions

- lacksquare $\mu(m{ heta}) := \mathbb{E}(D_{\mathcal{GP}}(m{ heta}))$ and $\mathrm{v}(m{ heta}) := \mathrm{Var}(D_{\mathcal{GP}}(m{ heta}))$
- ▶ Bayesian acquisition functions $A(\theta)$, quantify how 'desirable' it is to sample $\mathcal{D}(\theta)$ at θ .





Bayesian Acquisition Functions

- $lackbox{} \mu(m{ heta}) := \mathbb{E}(D_{\mathcal{GP}}(m{ heta})) \text{ and } \mathrm{v}(m{ heta}) := \mathrm{Var}(D_{\mathcal{GP}}(m{ heta}))$
- ▶ Bayesian acquisition functions $A(\theta)$, quantify how 'desirable' it is to sample $\mathcal{D}(\theta)$ at θ .
- Gutmann and Cor 2016 minimise lower confidence bound

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

the lower value of a confidence interval (determined by η_t).





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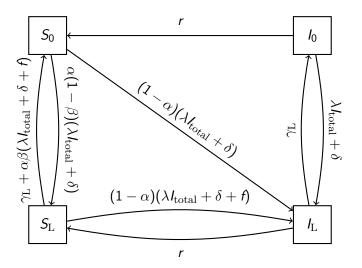




Figure: Champagne et al. 2022 P. vivax model



Champagne Model Parameters

- ightharpoonup lpha : proportion of those infected but cleared of blood stage infections (through treatment)
- β : 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
- $ightharpoonup \gamma_{\it L}$: rate of clearance of liver stage disease
- f: rate of relapse
- r: rate of blood stage clearance





Model Simulation

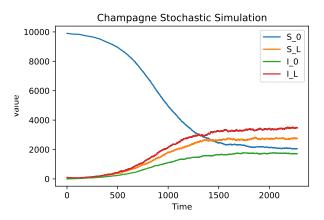


Figure: Exact stochastic simulation using parameters reported in Champagne et al. 2022. Population 10,000, initial infections 100.





'Observed' Data

- $ightharpoonup \mathbf{Y}_{\mathsf{obs}} := \{\iota_{\mathsf{obs}}, \pi_{\mathsf{obs}}, i_{\mathsf{obs}}, p_{\mathsf{obs}}\}$
 - $ightharpoonup \iota_{
 m obs}$: weekly incidence around steady state equilibrium
 - \blacktriangleright π_{obs} : prevalence around steady state equilibrium
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 - \triangleright p_{obs} : prevalence after one month of simulation
- \triangleright $\mathcal{D}(\alpha, \beta, \gamma_L, \lambda, f, r)$ is the L_2 norm of the relative differences

$$\sqrt{\left(\frac{\iota - \iota_{\text{obs}}}{\iota_{\text{obs}}}\right)^2 + \left(\frac{\pi - \pi_{\text{obs}}}{\pi_{\text{obs}}}\right)^2 + \left(\frac{i - i_{\text{obs}}}{i_{\text{obs}}}\right)^2 + \left(\frac{p - p_{\text{obs}}}{p_{\text{obs}}}\right)^2}$$





GP choices

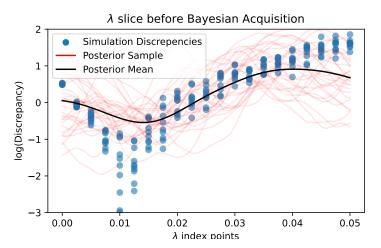
- \triangleright \mathcal{GP} choices
 - ightharpoonup Modelled In $\mathcal D$ as a Gaussian process
 - Matern kernel with $\nu = 5/2$
 - $\ell, \sigma_k^2, \sigma_o^2$ selected by leave one out cross validation.





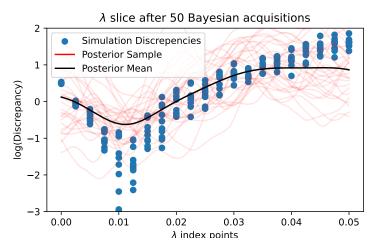






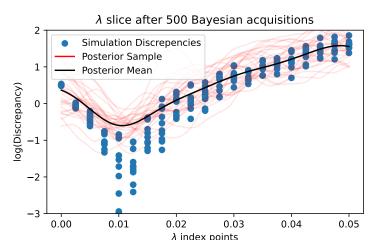






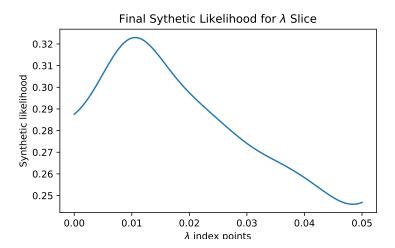






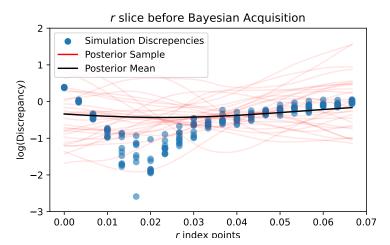






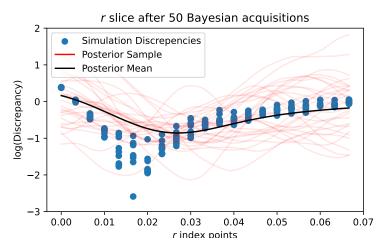






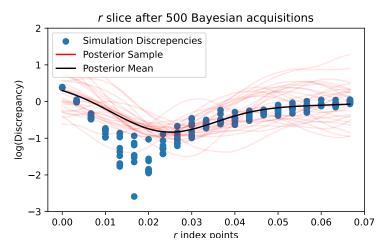






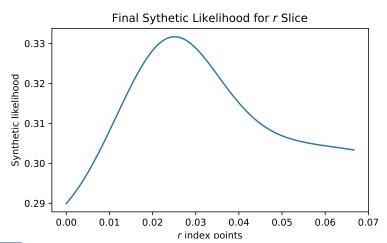
















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 - 1. observation variance,
 - 2. distribution of observations,
 - 3. behaviour of the discrepency mean.
- Possible extensions:





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- Possible extensions:
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 - 2. choose a different distribution and moment match,
 - 3. use a Student's *t*-process.





Conclusion

- Calibrating model parameters is important for scenario testing etc
- Successfully calibrated model parameters
- Could be used with more complicated models, even your model...





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- ► Eamon Conway
- ▶ Jennifer Flegg
- Ivo Mueller
- ► Mueller lab and unimelb MMB group





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