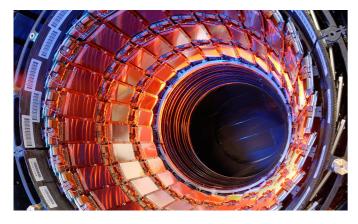
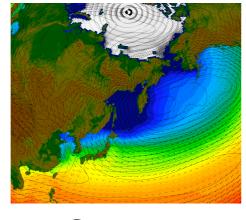
Scalable, reliably accurate Bayesian inference via approximate likelihoods and random features

Jonathan Huggins
Harvard University









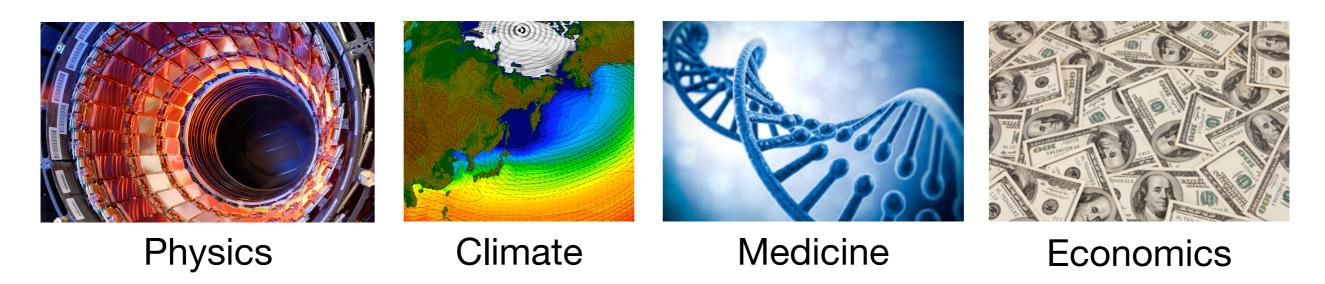
Physics

Climate

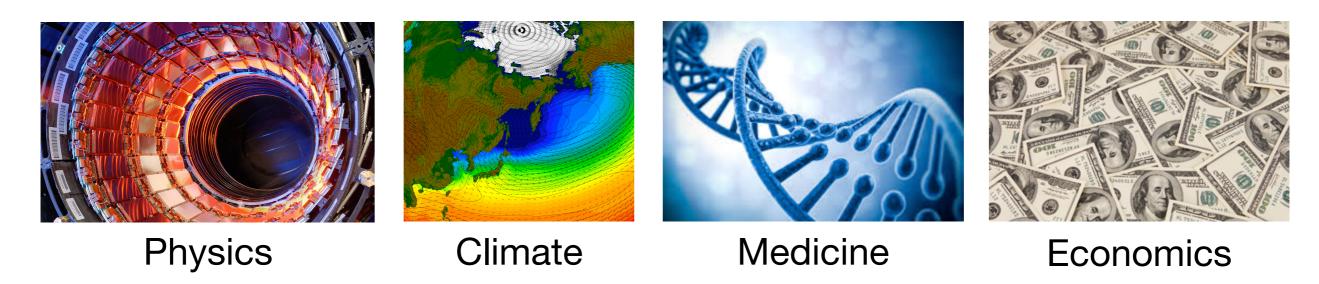
Medicine

Economics

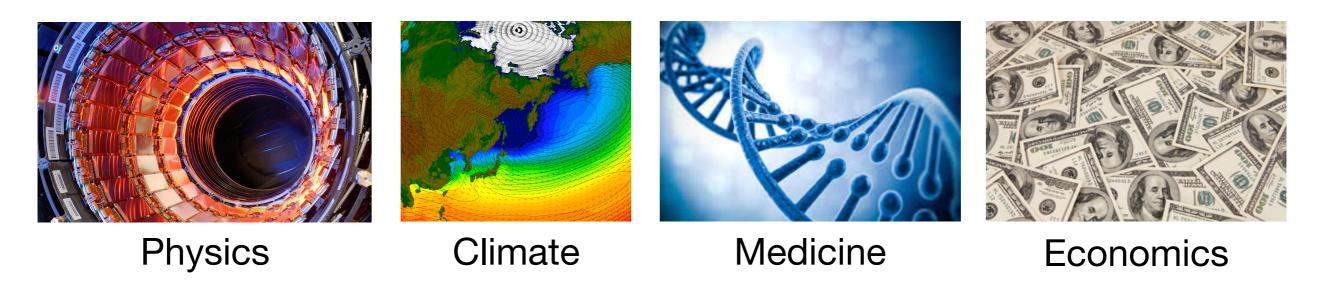
Large-scale data analysis for high-impact decision-making is widespread



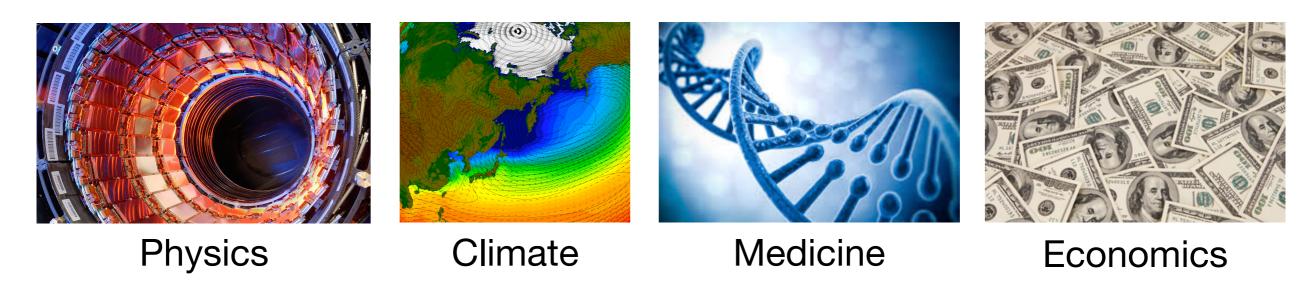
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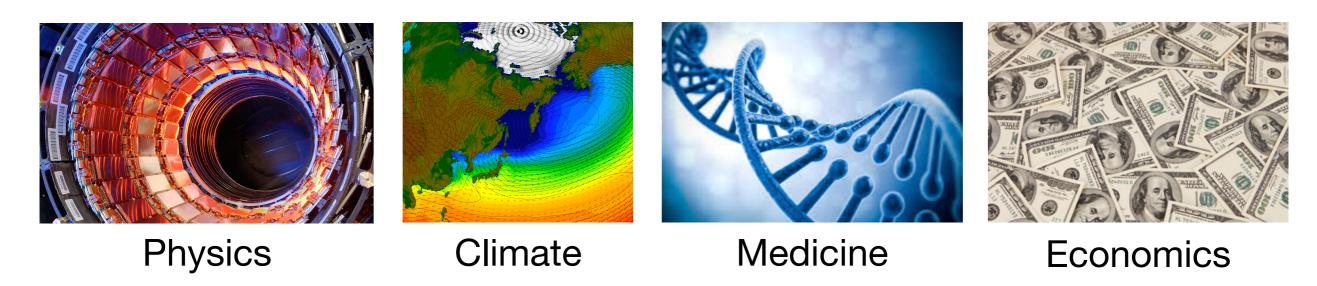
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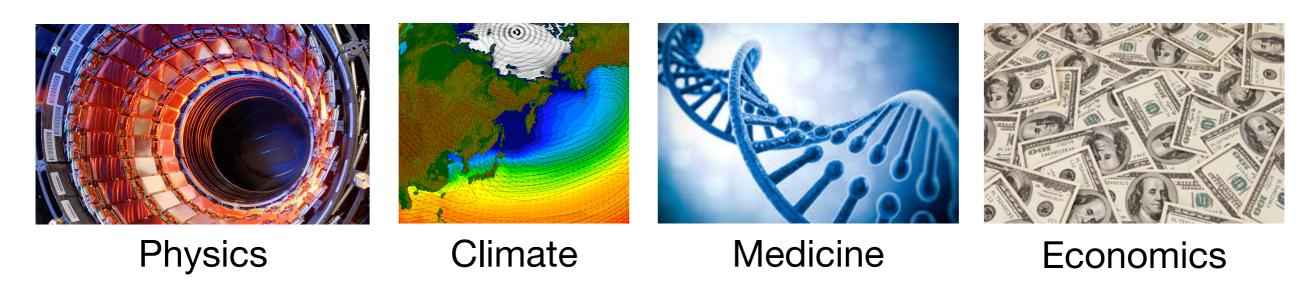
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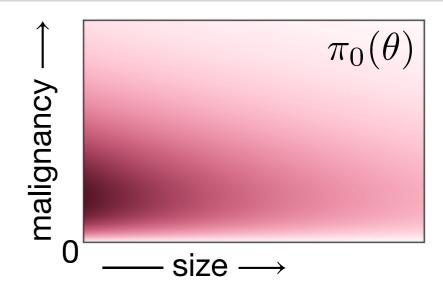
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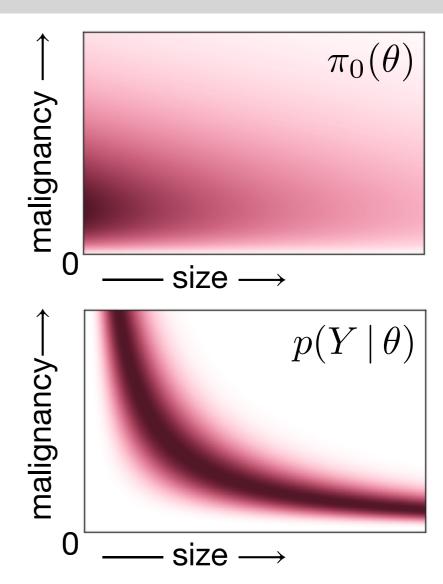
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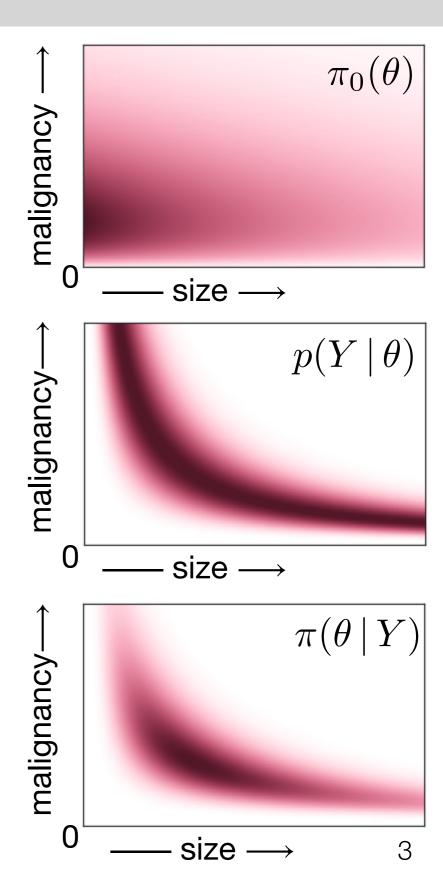


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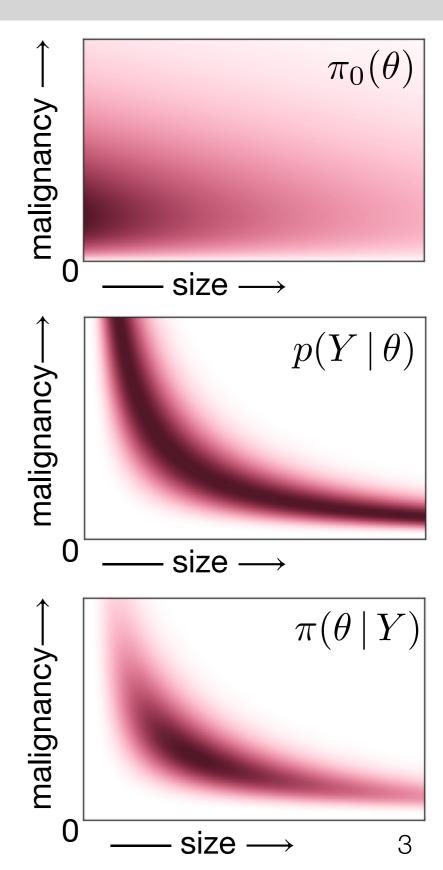
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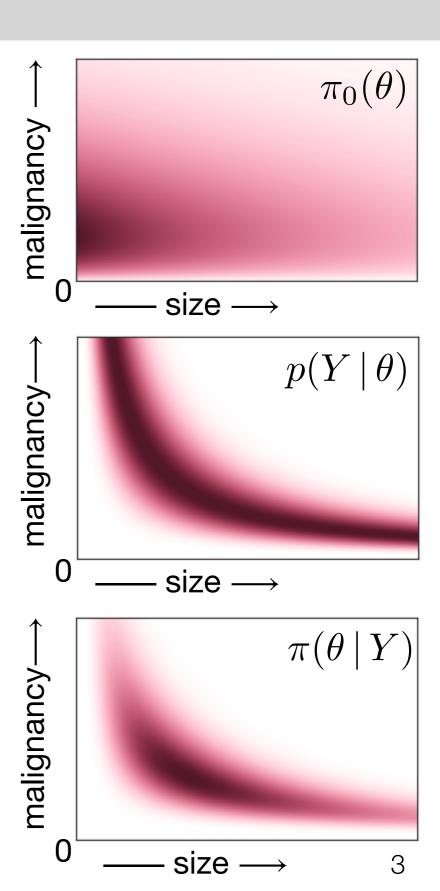


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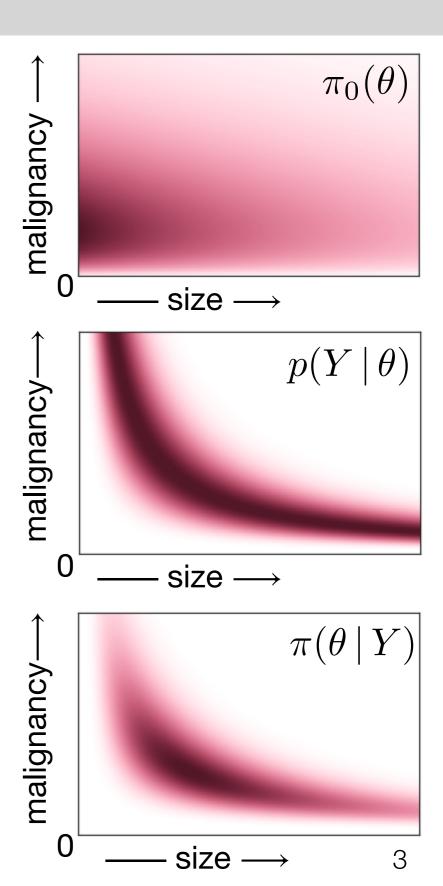
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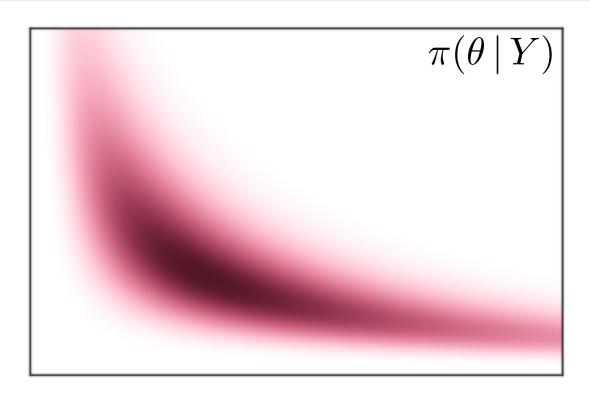
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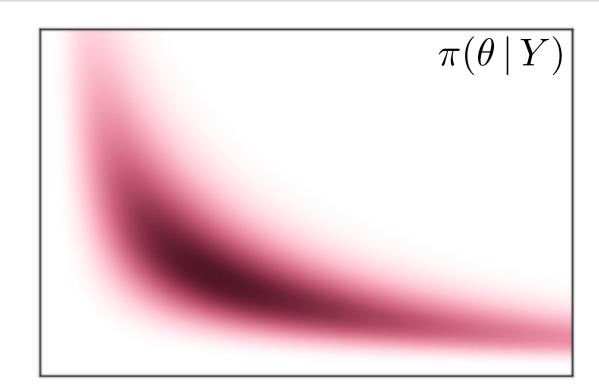
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 Computational challenges: posterior unnormalized, high-dimensional integral

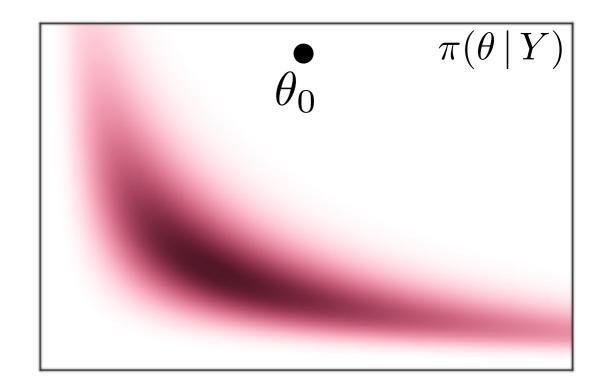




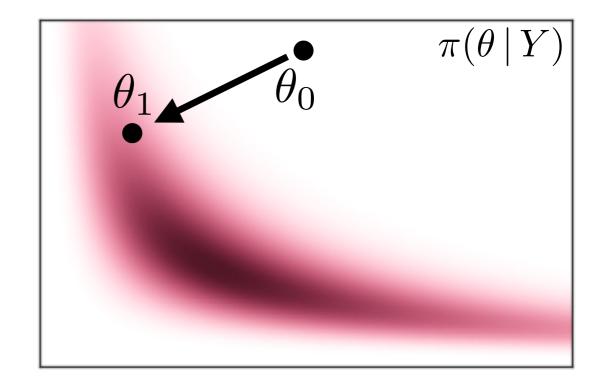
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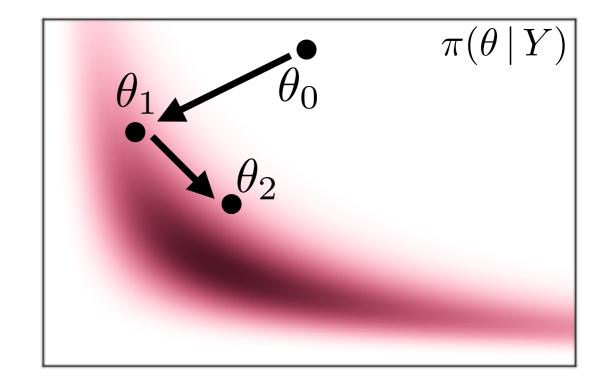
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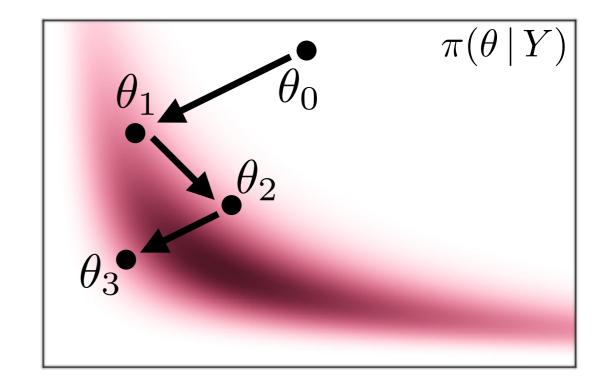
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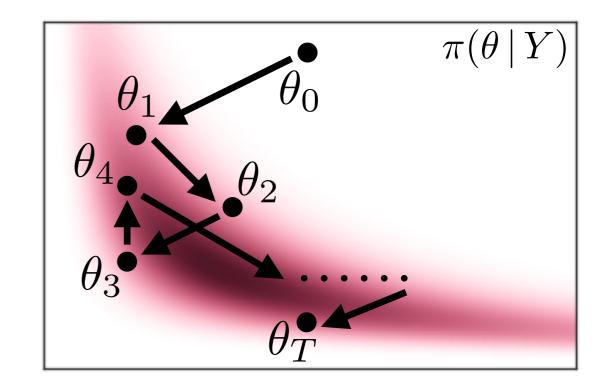
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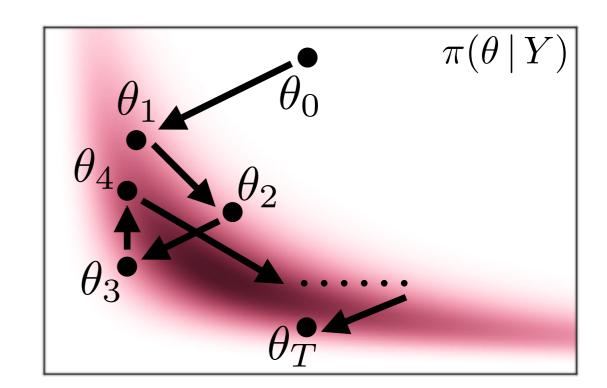


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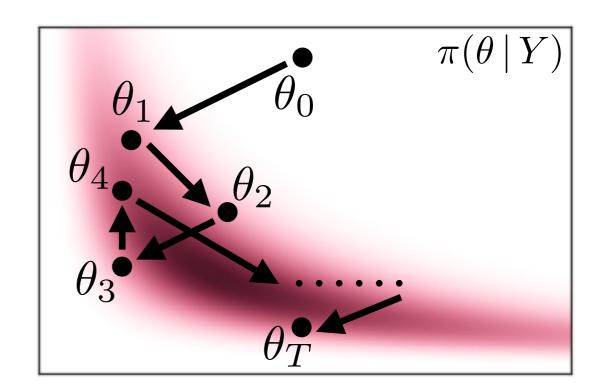
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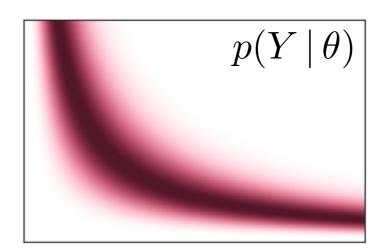


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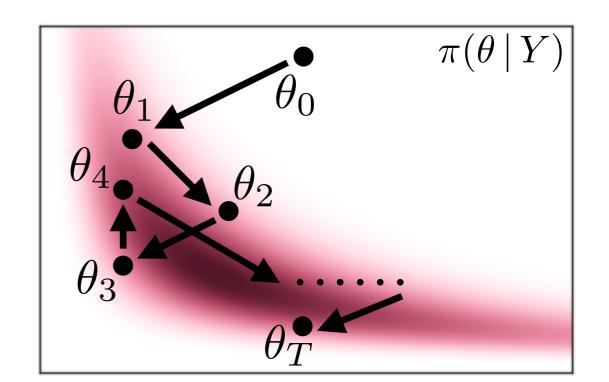


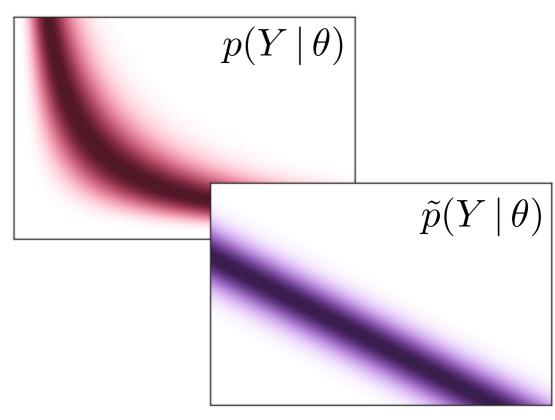


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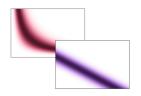
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- But MCMC is too slow: need to perform expensive evaluation of $p(Y | \theta_t)$ at iteration t
- Our scalable solution: use likelihood approximations that...
 - 1. Are accurate
 - 2. Are fast to compute
 - 3. Can be rigorously analyzed



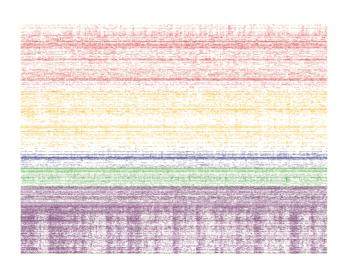


Agenda



A framework for scalable Bayesian inference

- Algorithm design
- Meaningful accuracy guarantees
- Validating results from heuristic algorithms

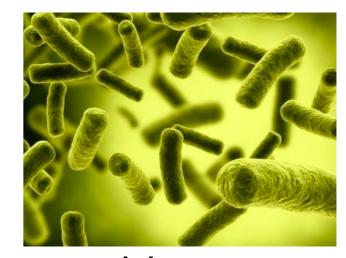


counts
[e.g. neural spikes]

Types of observations

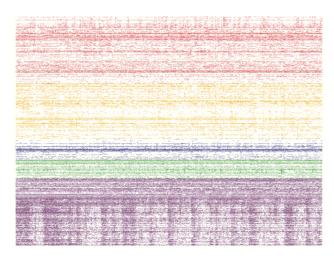


continuous [e.g. profit]



binary
[e.g. has disease?]

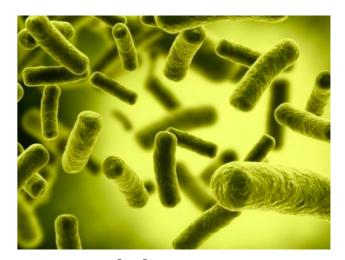




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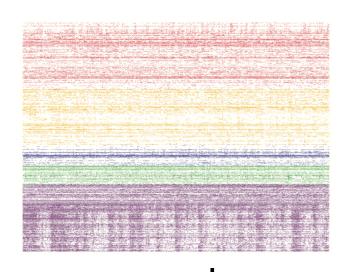


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Widely-adopted likelihood family: generalized linear models



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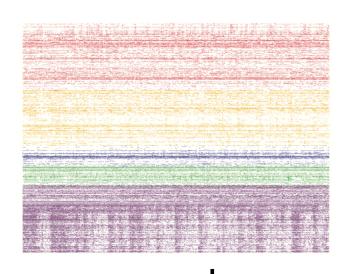
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Generalization of linear regression



counts
[e.g. neural spikes]

Types of observations



continuous [e.g. profit]



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Widely-adopted likelihood family: generalized linear models

- Generalization of linear regression
- Flexible, but still interpretable

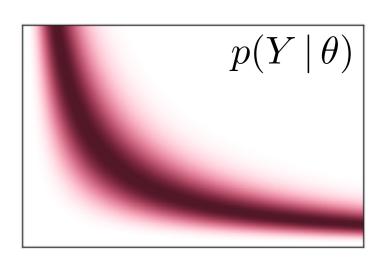
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Data $Y = \{y_1, y_2, \dots, y_N\}, y_n \in \mathbb{R}^d$, and parameter $\theta \in \mathbb{R}^p$

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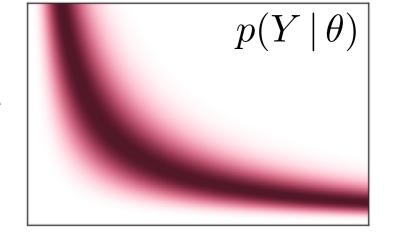
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We construct approximate sufficient statistics

Reparameterization function $\eta(\theta) \in \mathbb{R}^L$

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$$\eta(\theta) \in \mathbb{R}^{2}$$



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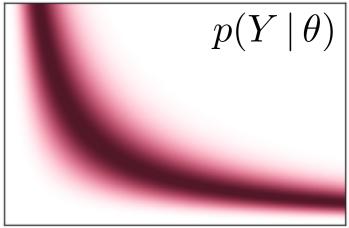
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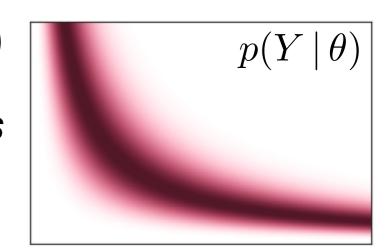
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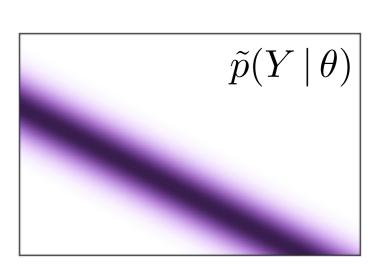
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Resulting approximation

$$\log p(Y \mid \theta) \approx \log \tilde{p}(Y \mid \theta) := \eta(\theta) \cdot \underbrace{\sum_{n=1}^{N} \tau(y_n)}_{\tau(Y)}$$



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Each component a polynomial: $\eta(\theta)_{\ell} \in \text{poly}(\theta), \tau(y_n)_{\ell} \in \text{poly}(y_n)$

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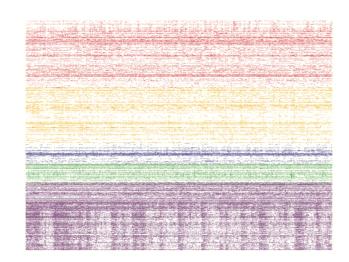
Polynomial approximations very accurate

counts
[e.g. neural spikes]

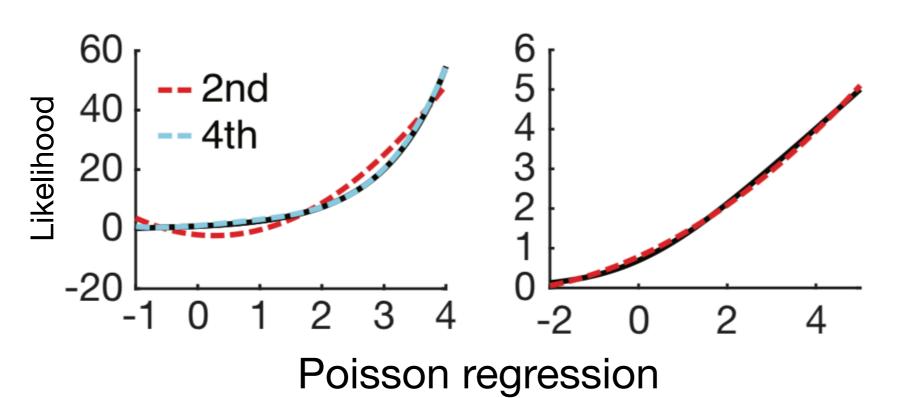


binary [e.g. has disease?]

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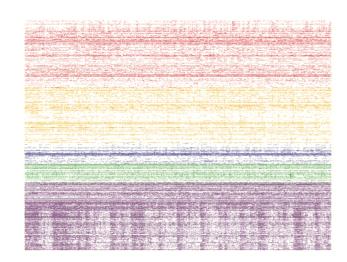


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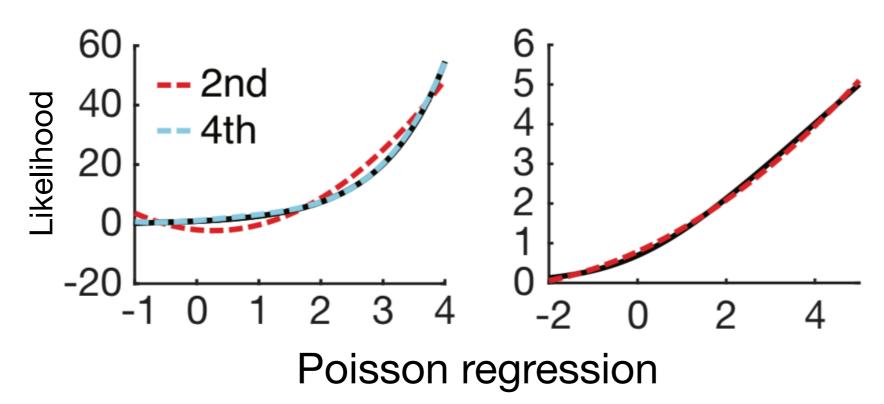


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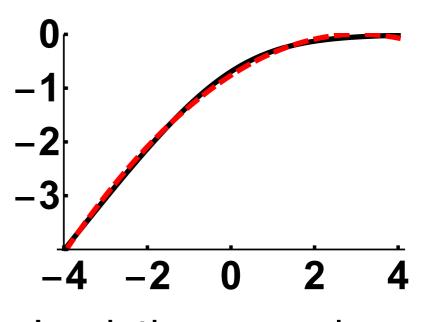
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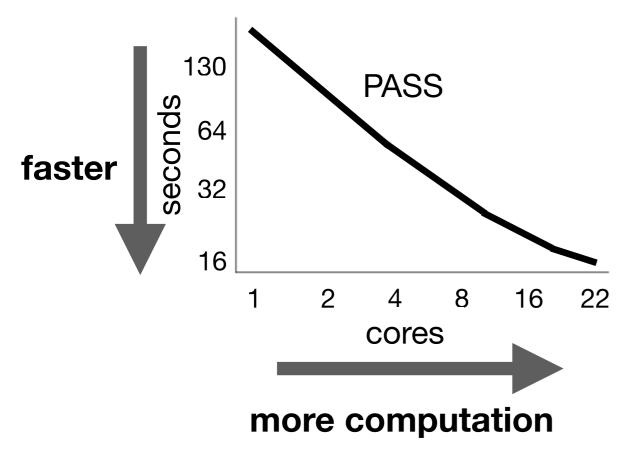
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Logistic regression

Fast, accurate empirical performance

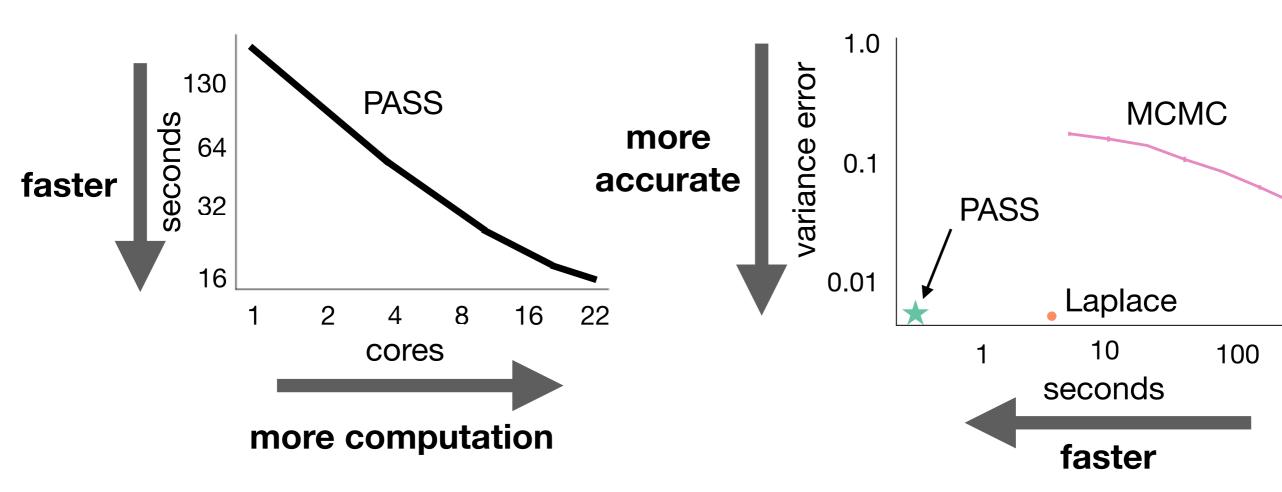
Fast distributed computation



- Logistic regression
- 6 million observations with 1,000 covariates
- MCMC: 1+ days

Fast, accurate empirical performance

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- Logistic regression
- 350,000 observations with 127 covariates

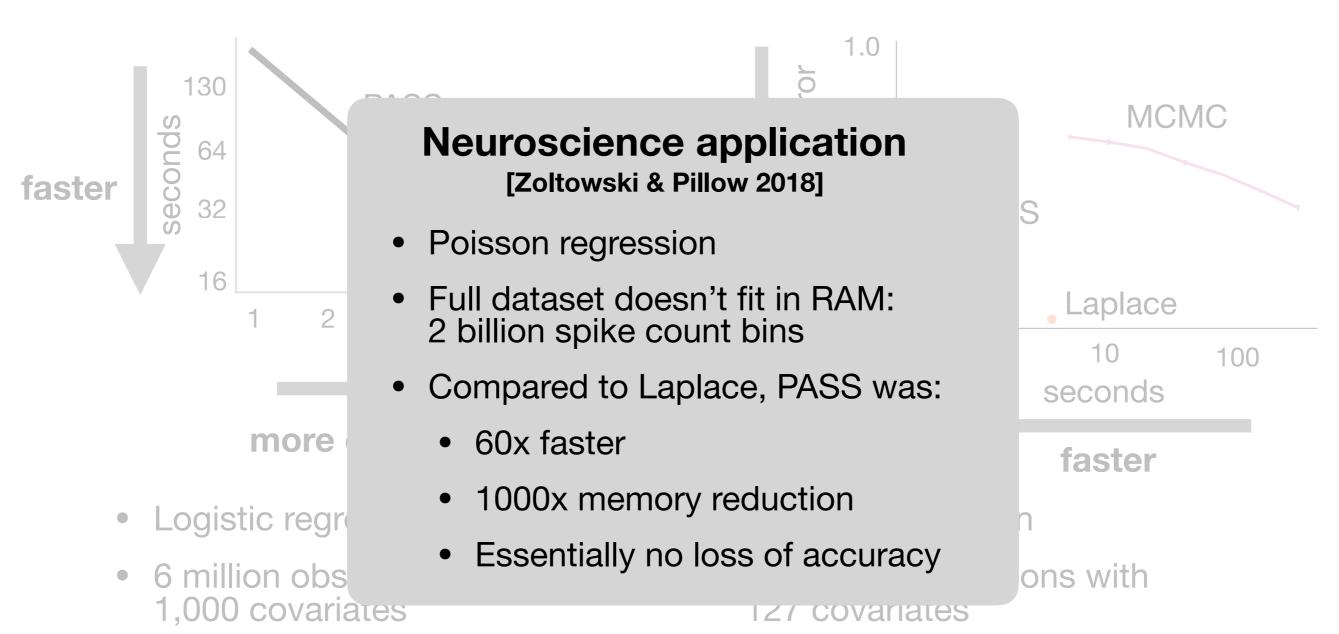
Fast and accurate

 Good mean estimation and predictive performance too

Fast, accurate empirical performance



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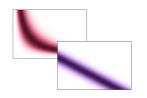
References

Huggins, Campbell, Kasprzak & Broderick. Scalable Gaussian process inference with finite-data mean and variance guarantees. AISTATS, 2019.

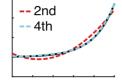
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Huggins, Campbell & Broderick. Coresets for scalable Bayesian logistic regression. Neural Information Processing Systems, 2016.

Agenda



A framework for scalable Bayesian inference



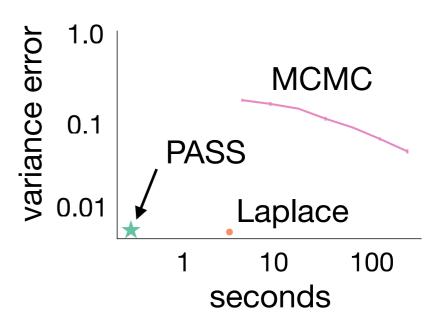
Algorithm design



Validating results from heuristic algorithms

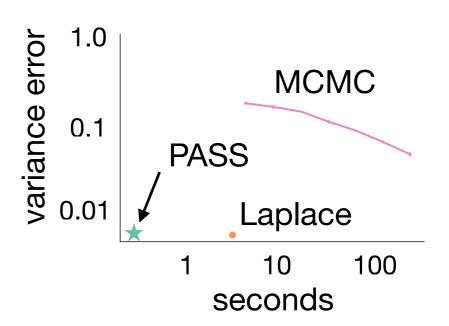
What about the next dataset?

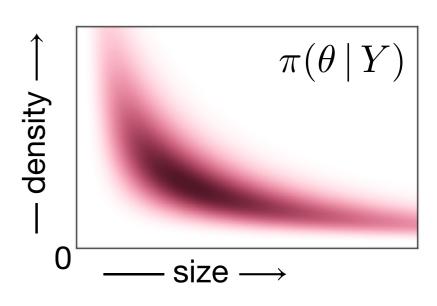
- Goal: Can we prove that PASS (or another likelihood approximation) will be accurate?
- If not, unsure if method is reliable



What about the next dataset?

- Goal: Can we prove that PASS (or another likelihood approximation) will be accurate?
- If not, unsure if method is reliable
- What's useful notion of accuracy?
- What do we want from the approximation?
 - Point estimate: mean
 - Uncertainty: standard deviation





Goal: good mean and standard deviation estimates

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- Computationally convenient: Kullback–Leibler divergence

$$\mathsf{KL}(q||\pi) = \mathbb{E}_q \left[\log \frac{q(\theta)}{\pi(\theta \mid Y)} \right] = \mathbb{E}_q \left[\log \frac{q(\theta)}{p(Y \mid \theta)\pi_0(\theta)} \right] + \text{constant}$$

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Proposition [HKCB18]

There exist q and π such that stdev(q)=1 and $stdev(\pi)=\infty$ but

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Proposition [HKCB18]

For Gaussians q and π such that stdev(q) = 1, it is possible that

$$|\operatorname{mean}(q) - \operatorname{mean}(\pi)| = e^{\mathsf{KL}(q||\pi)}$$

Better approximation properties: Wasserstein distance

$$W(\pi, q)^{2} = \inf_{\gamma \in \Gamma(\pi, q)} \int \|\theta - \theta'\|_{2}^{2} \gamma(d\theta, d\theta')$$

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- Goal: computational efficiency of Kullback–Leibler divergence and guarantees of Wasserstein distance

Many Wasserstein guarantees for MCMC...

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$$\nabla \log q(\theta)$$

Many Wasserstein guarantees for MCMC...

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$$\eta$$
-Fisher distance: $\mathsf{F}_{\eta}(\pi,q) = \mathbb{E}_{\eta} \left[\|\nabla \log \pi - \nabla \log q\|_2^2 \right]^{1/2}$

Many Wasserstein guarantees for MCMC...

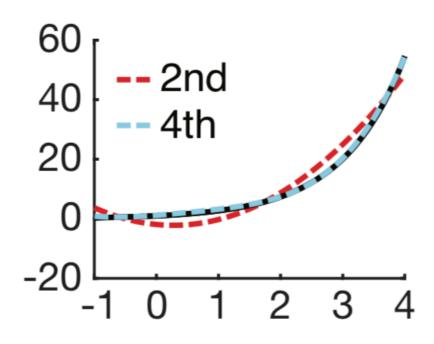
...but not for likelihood approximations

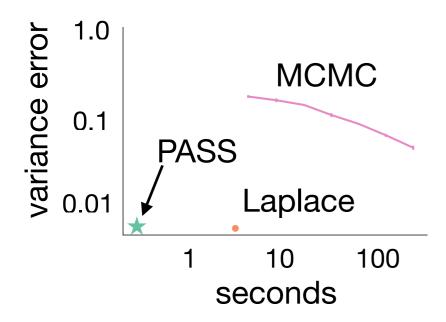
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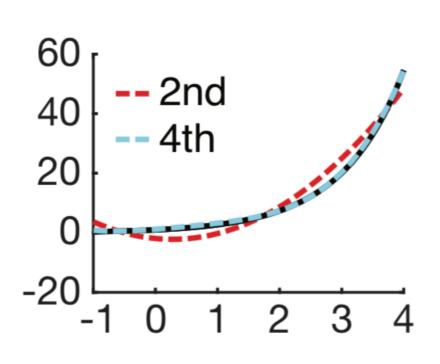
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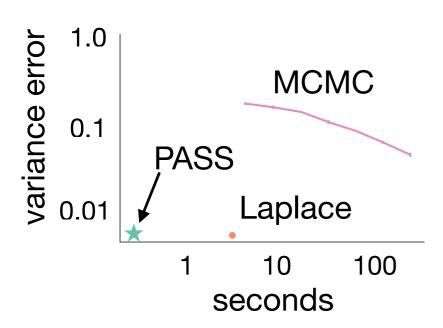
Theorem [HZ17, HKCB18]

$$W(\pi, q) \le C(q)C'(\eta, \pi)F_{\eta}(\pi, q)$$







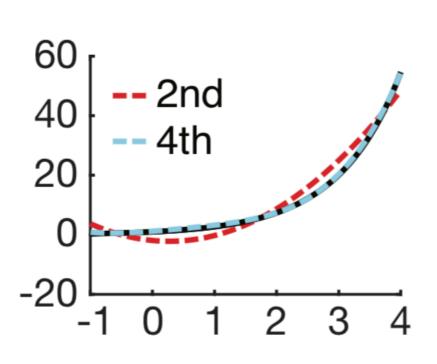


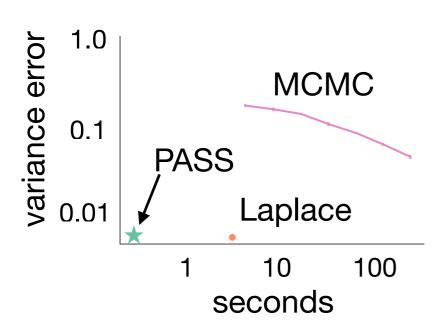
Theorem [HAB17]

Let q_M = the PASS approximate posterior using degree M polynomials.

Then the Wasserstein distance decreases exponentially in M:

$$W(\pi, q_M) \le cr^M$$





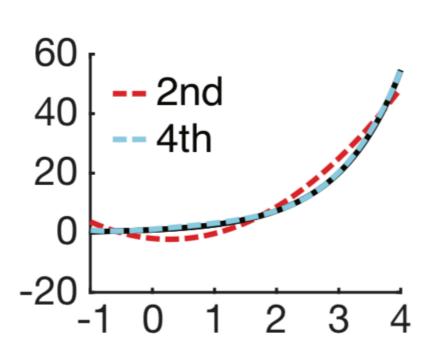
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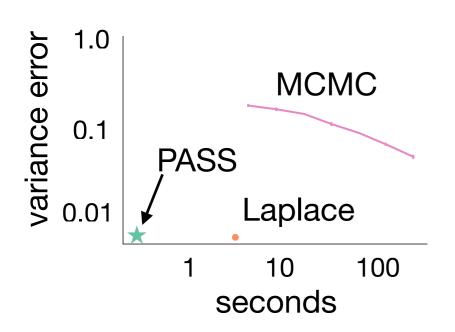
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Benefit: confidence to use PASS with a new dataset





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Then the Wasserstein distance decreases exponentially in M:

$$W(\pi, q_M) \le cr^M$$

- Benefit: confidence to use PASS with a new dataset
- Can also use η-Fisher distance to prove accuracy bounds for other likelihood approximations (e.g. Laplace approximation and coresets).

References

Theory

Huggins, Kasprzak, Campbell & Broderick. *Practical bounds on the error of Bayesian posterior approximations: A nonasymptotic approach.* arXiv:1809.09505 [stat.TH], 2018.

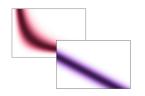
Huggins* & Zou*. Quantifying the accuracy of approximate diffusions and Markov chains. AISTATS, 2017.

Applications

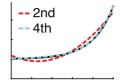
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Agenda



A framework for scalable Bayesian inference

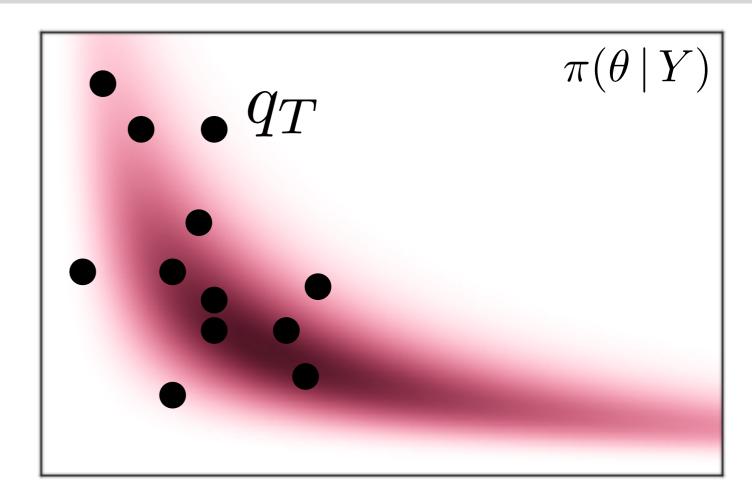


Algorithm design

F_n Meaningful accuracy guarantees



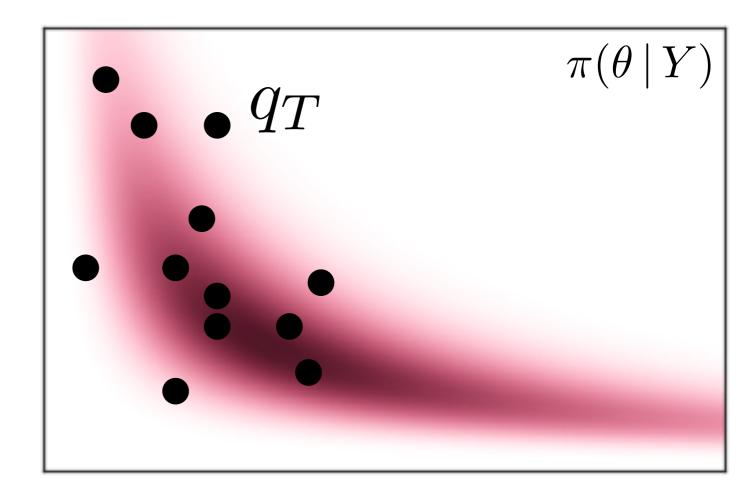
Validating results from heuristic algorithms



Goals

1. approximation quality

$$q_T \approx \pi$$
?



Goals

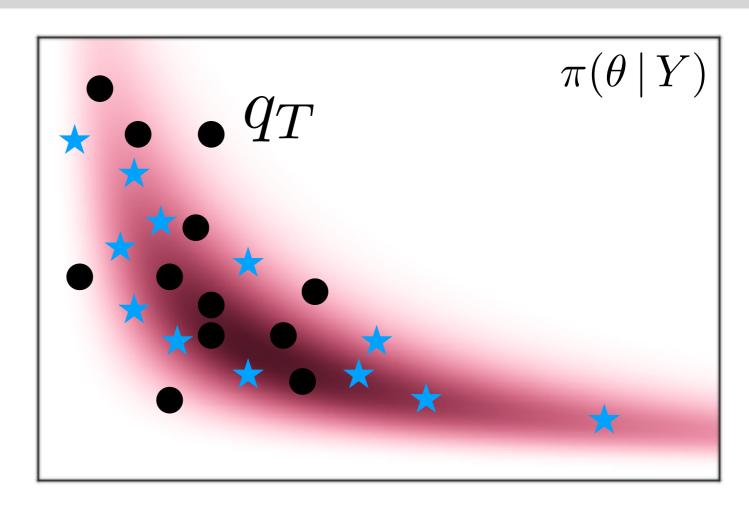
1. approximation quality $q_T \approx \pi$?

2. algorithm selection



versus





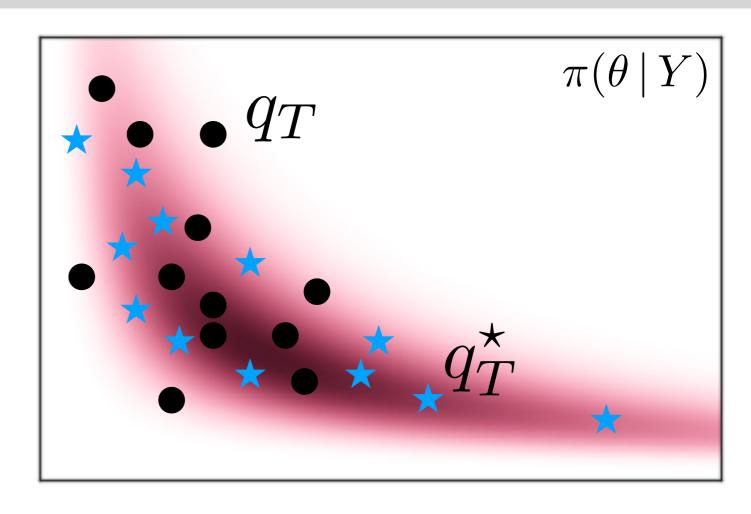
Goals

- **1.** approximation quality $q_T \approx \pi$?
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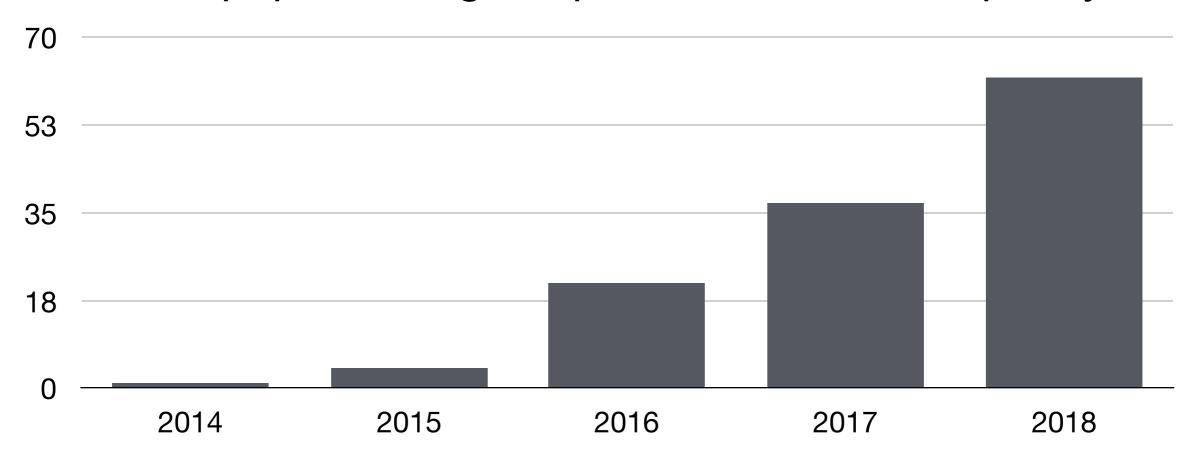


Approach: use a discrepancy measure $d(\pi, q_T)$

- Goal 1: is $d(\pi, q_T) \approx 0$?
- Goal 2: is $d(\pi, q_T^{\star})$ or $d(\pi, q_T)$ smaller?

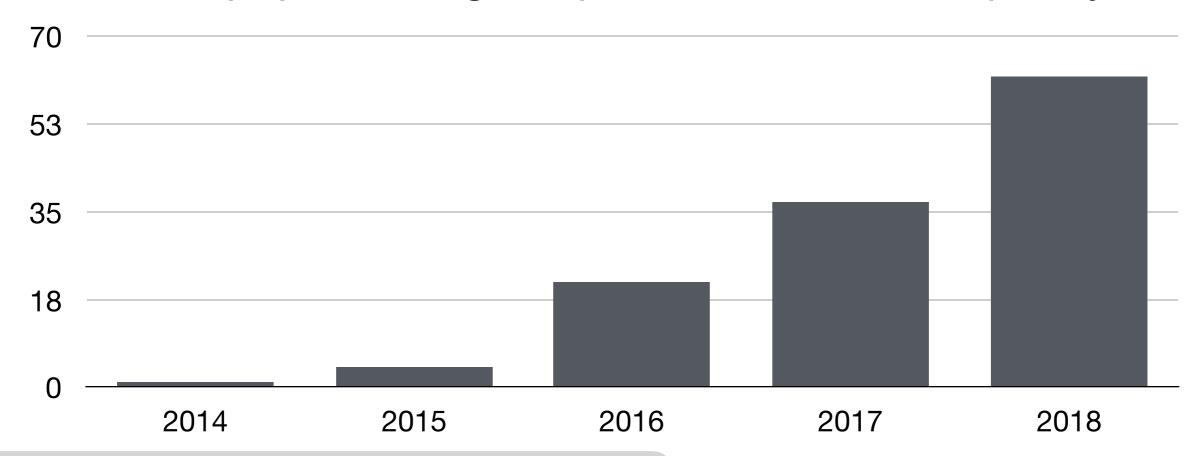
Approach: Stein discrepancies

of papers using the phrase "Stein discrepancy"



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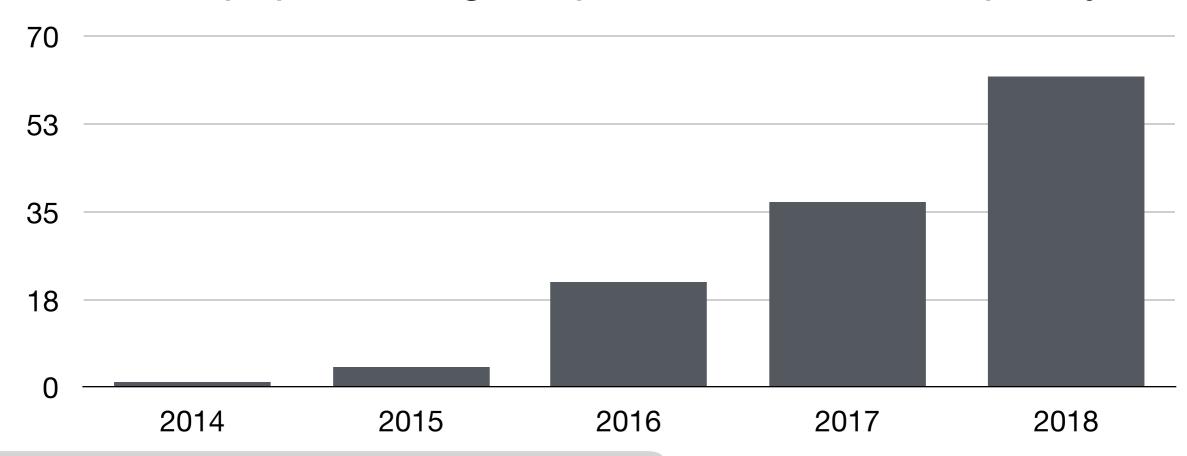


Definition

 $d(\pi, q_T)$ is theoretically sound if it detects (non-)convergence of $q_T \to \pi$ as $T \to \infty$

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Definition

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We provide the first discrepancy measure that is

- √ fast
- ✓ theoretically sound

$$q_T = T^{-1} \sum_{t=1}^T \delta_{\theta_t}$$
 kernel $k : \mathbb{R}^d \times \mathbb{R}^d \to \mathbb{R}$

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Theoretically sound approach: kernel Stein discrepancies (KSDs)

$$S_k(\pi, q_T) = T^{-2} \sum_{t=1}^T \sum_{t'=1}^T (\mathcal{T}_{\pi} \otimes \mathcal{T}_{\pi}) k(\theta_t, \theta_{t'})$$

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$$S_k(\pi, q_T) \approx M^{-1} \sum_{m=1}^M \rho(Z_m)^{-1} \left\{ T^{-1} \sum_{t=1}^T \mathcal{T}_{\pi} \psi(\theta_t, Z_m) \right\}^2$$

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• $\Theta(MT)$ time complexity but not theoretically sound

Our solution: design Stein discrepancy from the start for use with importance sampling

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Feature Stein discrepancies are theoretically sound:

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Recall: $\Theta(MT)$ time complexity when using M importance samples

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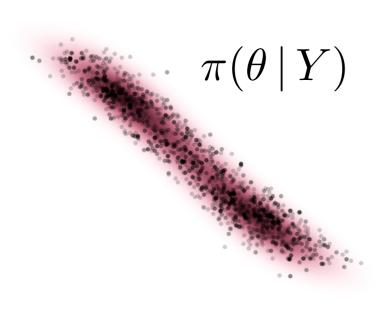
Theorem [HM18]

For any $\alpha>0$, we can compute a theoretically sound random feature Stein discrepancy using $M=\Theta(T^{\alpha})$ importance samples in near-linear $\Theta(T^{1+\alpha})$ time.

"exact" MCMC

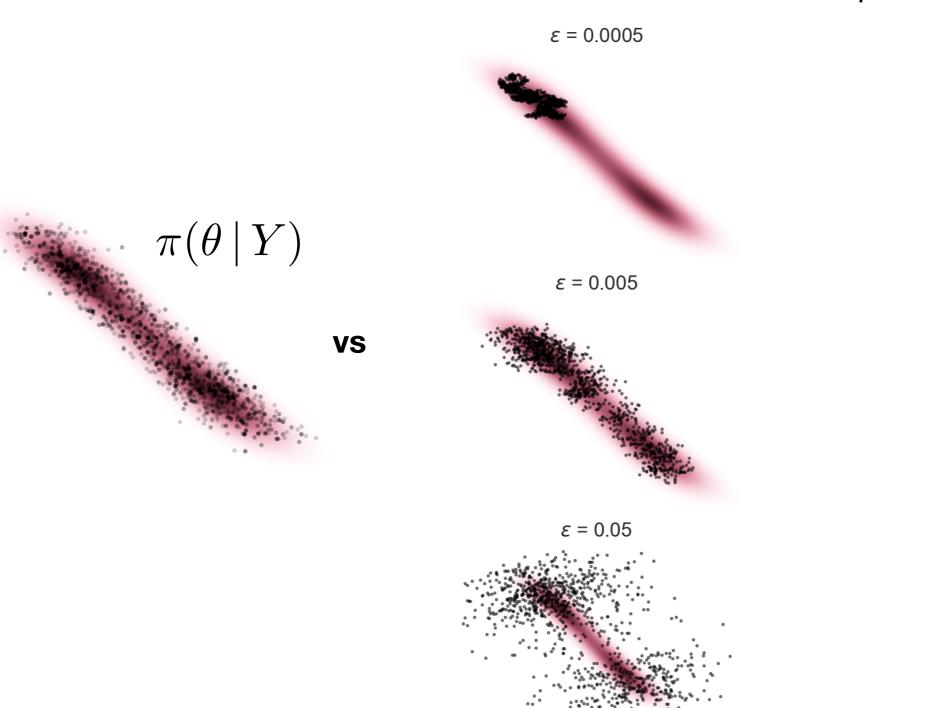
$$\pi(\theta \mid Y)$$

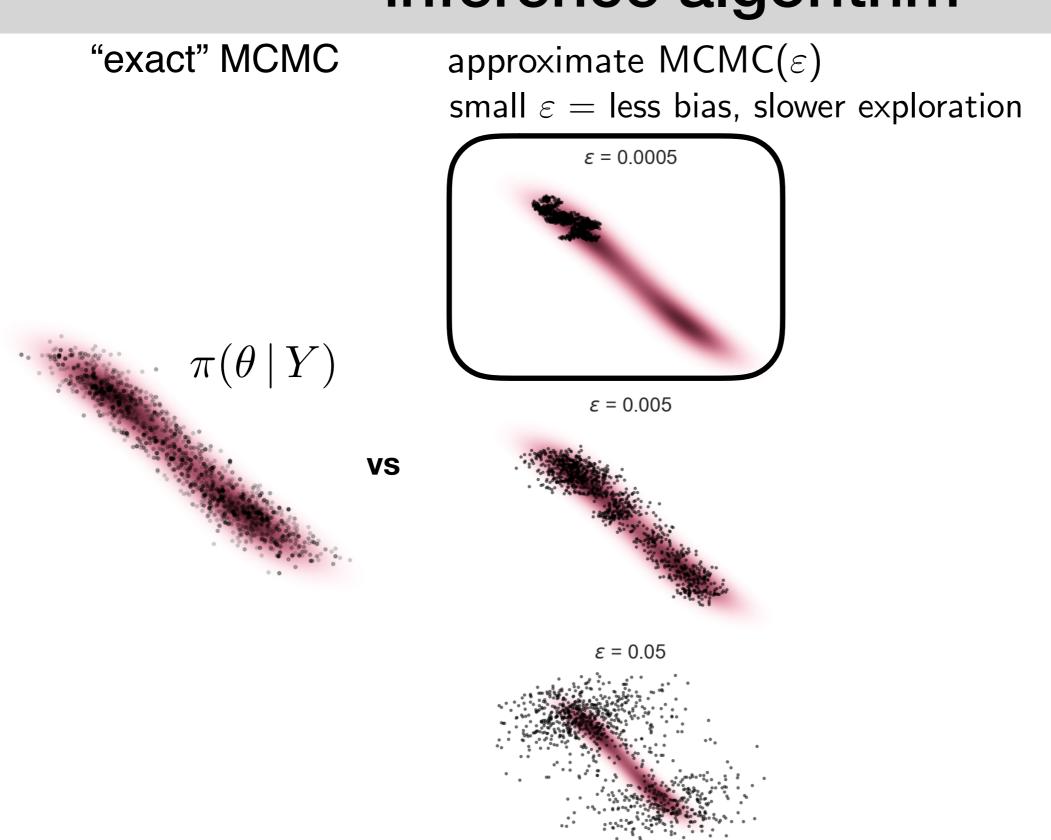
"exact" MCMC



"exact" MCMC

approximate $MCMC(\varepsilon)$ small ε = less bias, slower exploration

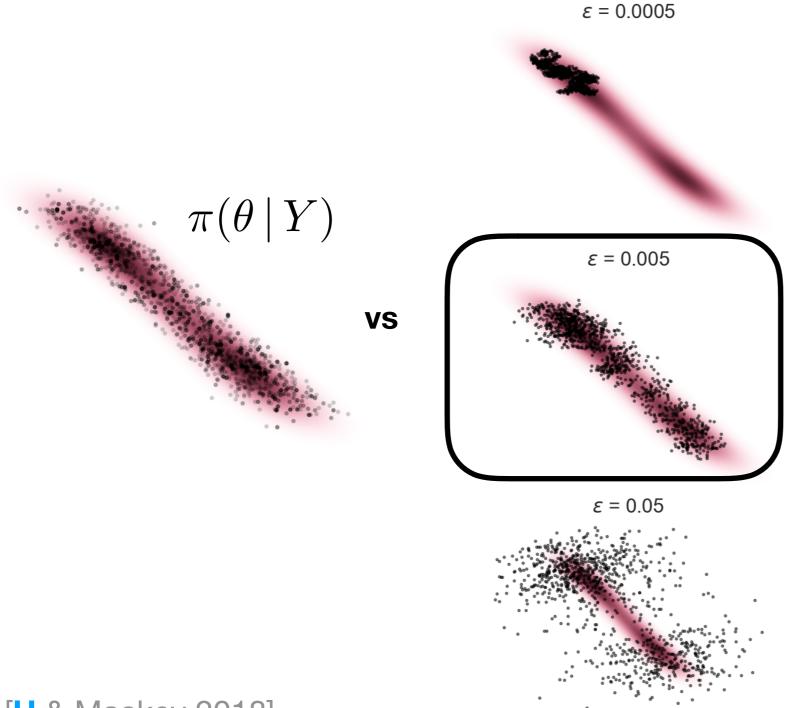




"exact" MCMC approximate $MCMC(\varepsilon)$ small $\varepsilon = less bias$, slower exploration $\varepsilon = 0.0005$ $\pi(\theta \mid Y)$ $\varepsilon = 0.005$ **VS** $\varepsilon = 0.05$

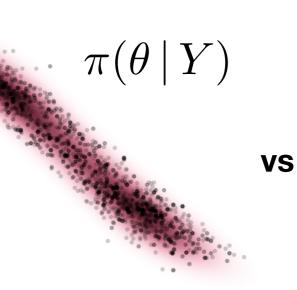
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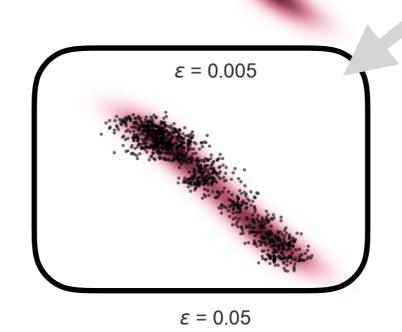
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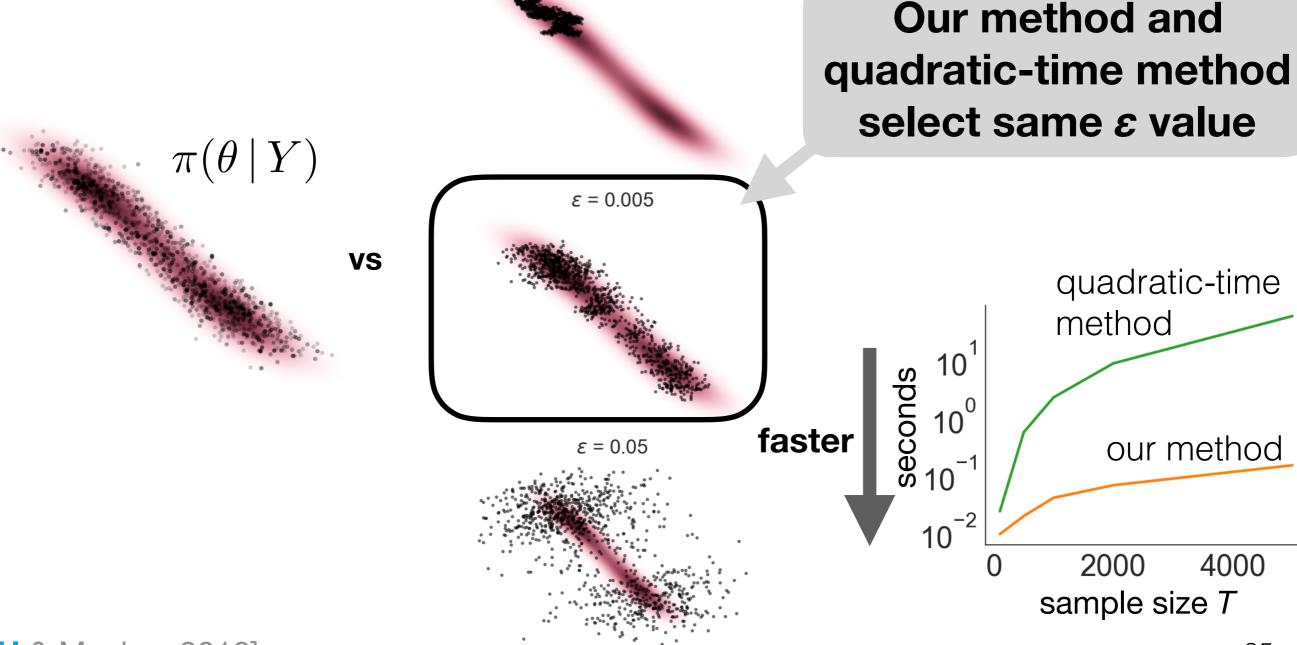




Our method and quadratic-time method select same ε value

"exact" MCMC

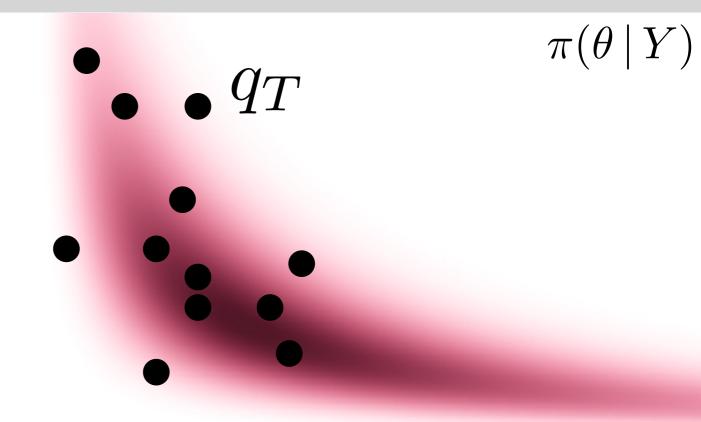
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[H & Mackey 2018]

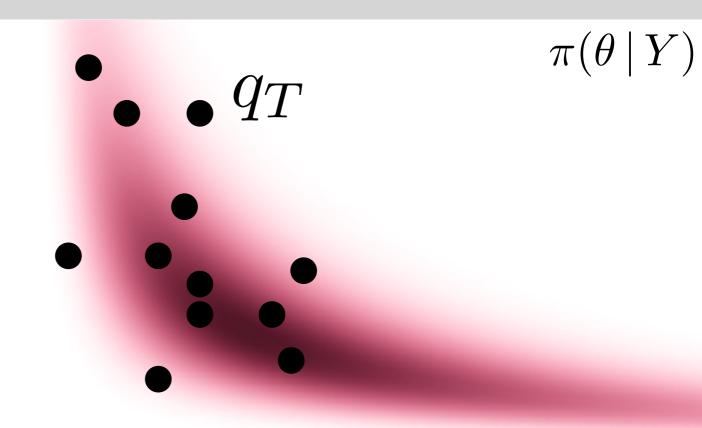
Application #2: goodness-of-fit testing

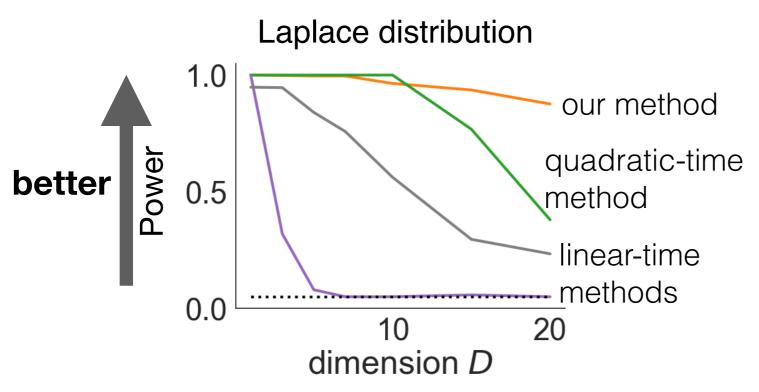
- Question: $q_T \approx \pi$?
- Power = probability of correctly rejecting null
- π = standard Gaussian

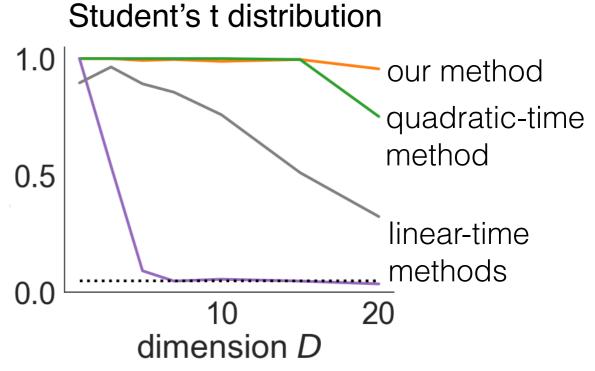


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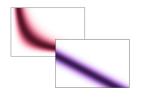
[H & Mackey 2018]

References

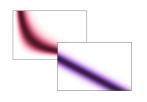
Agrawal, Campbell, Huggins & Broderick. Data-dependent compression of random features for large-scale kernel approximation. AISTATS, 2019.

Huggins* & Mackey*. Random feature Stein discrepancies. Neural Information Processing Systems, 2018.

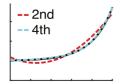
Huggins, Kasprzak, Campbell & Broderick. *Practical bounds on the error of Bayesian posterior approximations: A nonasymptotic approach.* arXiv:1809.09505 [stat.TH], 2018.



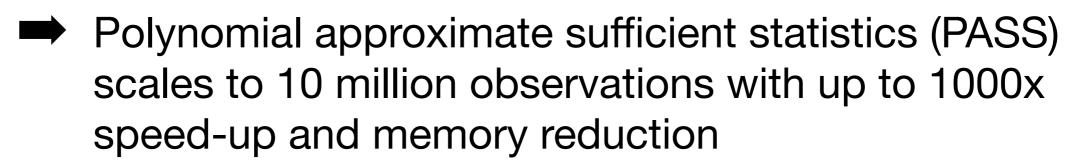
A framework for scalable Bayesian inference

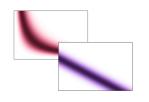


A framework for scalable Bayesian inference

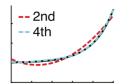


Algorithm design





A framework for scalable Bayesian inference

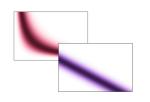


Algorithm design

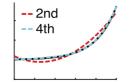
→ Polynomial approximate sufficient statistics (PASS) scales to 10 million observations with up to 1000x speed-up and memory reduction

F_η Meaningful accuracy guarantees

General theory for obtaining practical error bounds for likelihood approximations, including PASS



A framework for scalable Bayesian inference



Algorithm design

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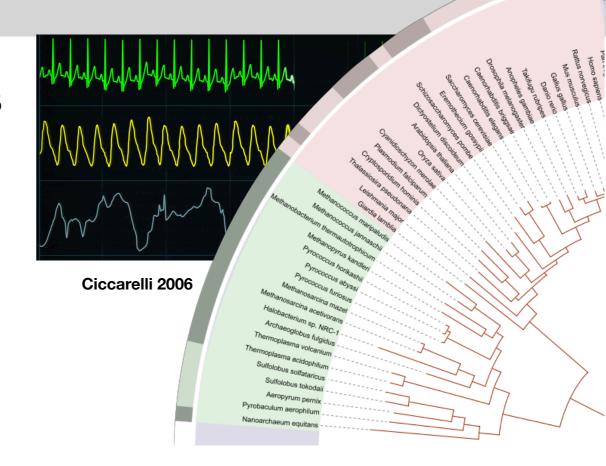
General theory for obtaining practical error bounds for likelihood approximations, including PASS

Validating results from heuristic algorithms

→ Fast (near-linear time) and theoretically sound Stein discrepancy measure

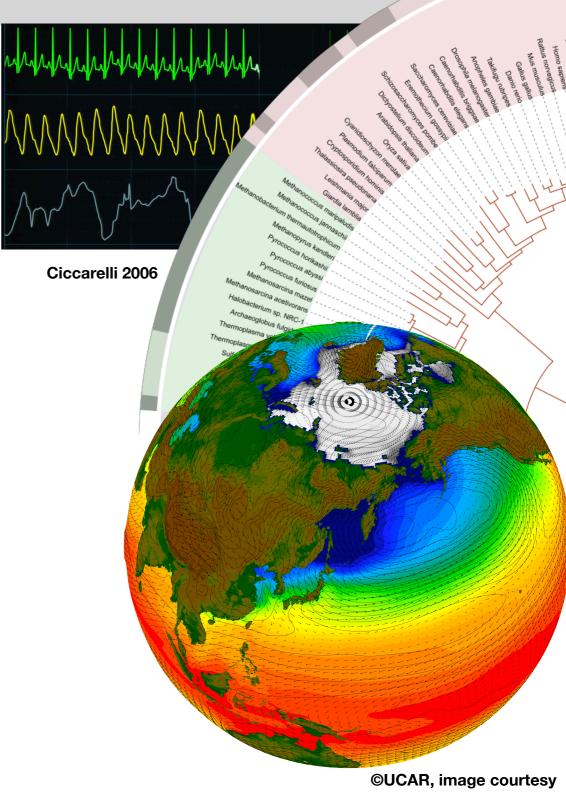
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 Scalable inference for time series and other structured data
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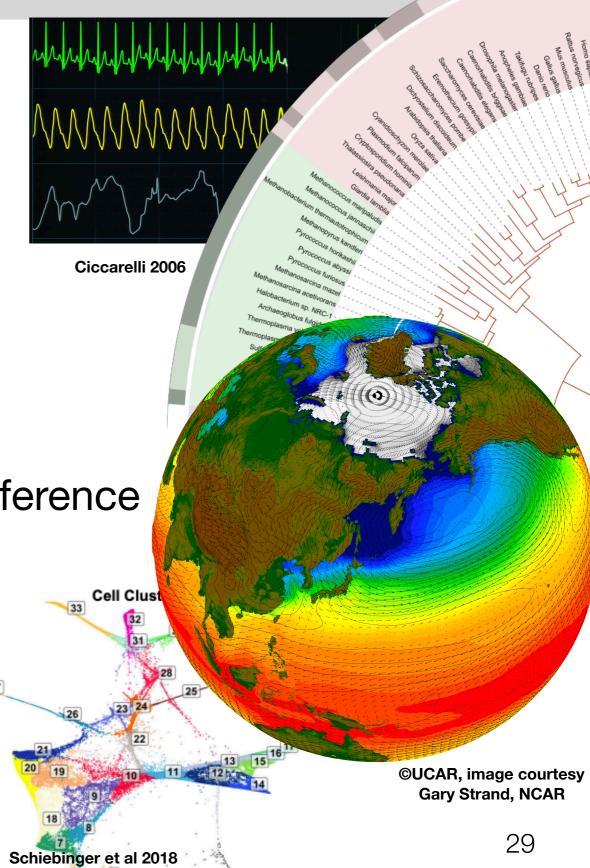
Future Directions

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 [e.g. phylogenetic trees]

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Statistically robust yet scalable inference
 [e.g. in cancer genomics]





References

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Markov chain Monte Carlo

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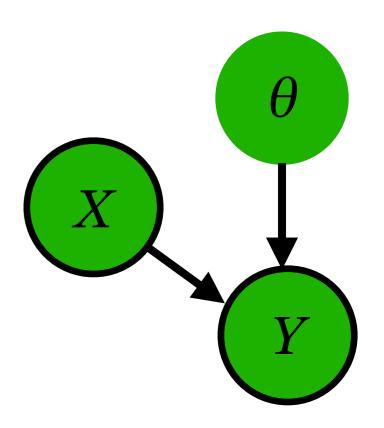
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Theory provides loose quantitative rates	e.g. geometric ergodicity	e.g. linear or 1/T	e.g. exponentially small error
Theory does not apply to many real-world cases	e.g. non-trivial hierarchical models	e.g. non-convex functions	e.g. non-trivial hierarchical models
Practical methods to evaluate success	e.g. Gelman–Rubin diagnostic	e.g. norm of gradient, duality gap	e.g. Stein discrepancies
Comparison of algorithms	e.g. effective samples per second	e.g. time to convergence	e.g. Stein discrepancies

PASS for generalized linear models (PASS-GLM)

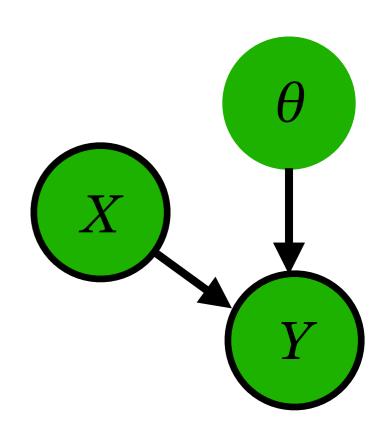
$$Y = \{y_1, y_2, \dots, y_N\}, y_n \in \mathbb{R}, X = \{x_1, x_2, \dots, x_N\}, x_n \in \mathbb{R}^d, \theta \in \mathbb{R}^d$$



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GLMs: $\log p(y_n \mid x_n, \theta) = \phi(y_n, \theta \cdot x_n) \approx \eta(\theta) \cdot \tau(y_n, x_n)$

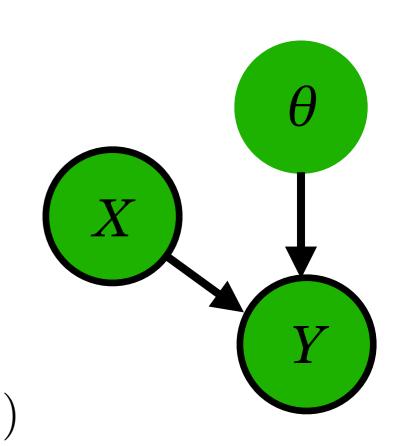


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$$\tau(y_n, x_n) = (y_n, x_{n1}, x_{n2}, \dots, x_{nd}, y_n^2, x_{n1}^2, x_{n2}^2, \dots, x_{nd}^2, y_n x_{n1}, \dots, y_n x_{nd}, x_{n1} x_{n2}, x_{n1} x_{n3}, \dots, y_n^M, x_{n1}^M, x_{n1}^M, x_{n2}^M, \dots, x_{nd}^M)$$



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$$L = \dim(\tau)$$

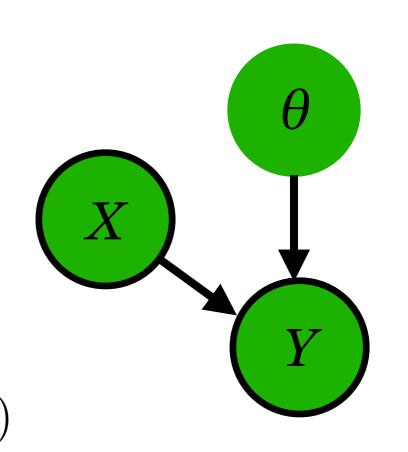
$$= {m+d+1 \choose m}$$

$$= O([d+1]^m)$$

$$= \dim(\tau) = \begin{pmatrix} y_n^2, x_{n1}^2, x_{n2}^2, \dots, x_{nd}^2, \\ y_n x_{n1}, \dots, y_n x_{nd}, \\ x_{n1} x_{n2}, x_{n1} x_{n3}, \dots, \\ \vdots \\ y_n^M, x_{n1}^M, x_{n2}^M, \dots, x_{nd}^M \end{pmatrix}$$

$$= 0([d+1]^m)$$

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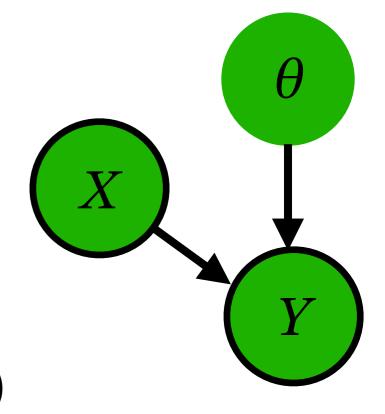
$$L = \dim(au)$$
 $y_n x_{n1}, \dots, y_n x_{nd},$ $= {m+d+1 \choose m}$ $x_{n1}x_{n2}, x_{n1}x_{n3}, \dots,$ $= O([d+1]^m)$ $\cdots,$

$$y_n^2, x_{n1}^2, x_{n2}^2, \dots, x_{nd}^2,$$

$$y_n x_{n1}, \ldots, y_n x_{nd},$$

$$x_{n1}x_{n2},x_{n1}x_{n3},\ldots,$$

$$y_n^M, x_{n1}^M, x_{n2}^M, \dots, x_{nd}^M$$



$$\tau(y_n, x_n) = \left(a(k, M) y_n^{k_0} \prod_{i=1}^d x_{ni}^{k_i} \right)_{\substack{k \in \mathbb{N}^{d+1} \\ \sum_i k \le M}} \eta(\theta) = \left(\prod_{i=1}^d \theta_i^{k_i} \right)_{\substack{k \in \mathbb{N}^{d+1} \\ \sum_i k \le M}}$$