

# Some applications of MCMC perturbations in high-dimensional problems

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# Introduction

- ▶ Aim: develop scalable MCMC algorithms for large  $(N, p)$  regression with continuous shrinkage priors
- ▶ Compute the posterior expectation & marginal posterior densities for the coefficients
- ▶ We won't get this from optimization, also not a convex problem in many cases
- ▶ For concreteness, we focus on the horseshoe prior of Carvalho et al. (2010) - theoretical support + empirical performance
- ▶ Basic ingredients extend to more general Gaussian variance mixtures as well as two-component mixtures like the spike-and-slab lasso (Rockova & George, 2014)

# Approximations in MCMC

- ▶ Our proposed algorithm introduces certain approximations at each MCMC step - approximate certain expensive matrix multiplications
- ▶ Leads to substantial computational advantages
- ▶ How to quantify the effect of such approximations?
- ▶ Perturbation theory for MCMC algorithms (Alquier et al. 2014, Rudolf & Schweizer (2018), Johndrow & Mattingley (2018)...)
- ▶ A new general result + bounds on approximation error for our algorithm

## Other applications

- ▶ Similar ideas applicable to a host of other high-dimensional problems
- ▶ Ongoing work: approximate sampling from truncated multivariate normals with applications to problems with constrained parameters
- ▶ Replace the hard constraints with “soft” versions

## Bayesian shrinkage: motivation and background

## “Global-local” shrinkage priors

- ▶ Consider a Gaussian linear model

$$z = W\beta + \varepsilon, \quad \varepsilon \sim N(0, \sigma^2 I_N)$$

where  $W$  is  $N \times p$ , with  $N, p$  both possibly large

- ▶ The basic form of the prior is

$$\beta_j \mid \sigma, \xi, \eta \stackrel{ind}{\sim} N(0, \sigma^2 \xi^{-1} \eta_j^{-1})$$

- ▶ The  $\eta_j^{-1/2}$  are the “local scales” and  $\xi^{-1/2}$  the “global scale”
- ▶ A popular choice for  $\pi(\xi, \eta)$  is the “Horseshoe” (Carvalho et al. 2010)

$$\eta_j^{-1/2} \stackrel{ind.}{\sim} \text{Cauchy}_+(0, 1), \quad \xi^{-1/2} \sim \text{Cauchy}_+(0, 1)$$

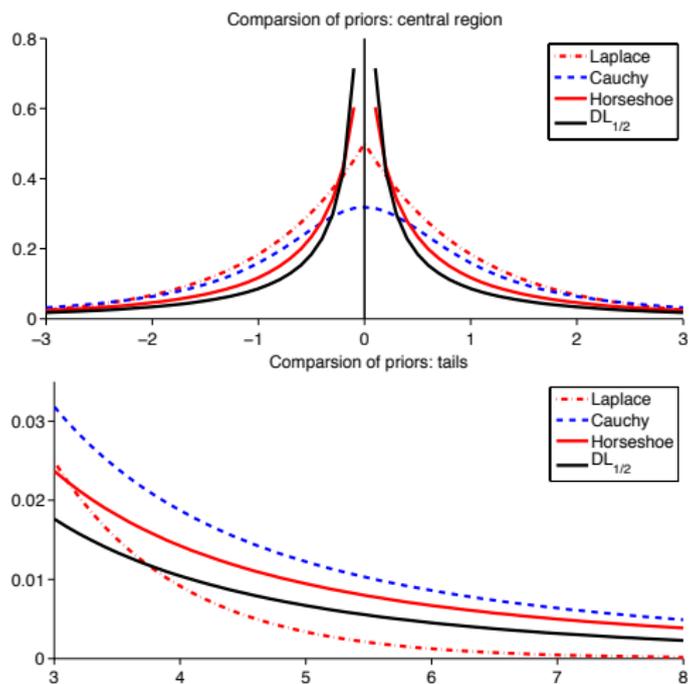
## “Global-local” shrinkage priors

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- ▶ The  $\eta_j^{-1/2}$  are the “local scales” and  $\xi^{-1/2}$  the “global scale”
- ▶ Only global scale  $\Rightarrow$  ridge type shrinkage
- ▶ Local scales help adapt to sparsity
- ▶ The global scale  $\xi^{-1/2}$  controls **how many**  $\beta_j$  are signals, and  $\eta_j^{-1/2}$  control **their identities**

# Continuous shrinkage via one group models



## Computational challenges

# MCMC review

- ▶ Basic idea of MCMC: construct a Markov transition kernel  $\mathcal{P}$  with invariant measure the posterior, i.e.  $\mu\mathcal{P} = \mu$  where  $\mu$  is the posterior measure
- ▶ Then approximate

$$\mu\varphi \equiv \int \varphi(x)\mu(dx) \approx n^{-1} \sum_{k=0}^{n-1} \varphi(X_k)$$

for  $X_k \sim \nu\mathcal{P}^{k-1}$

# Computational cost

- ▶ What is the computational cost? Two factors
  1. The cost of taking one step with  $\mathcal{P}$
  2. How long the Markov chain needs to be to make approximation “good”

## Computational cost per step

- ▶ Perform various matrix operations - multiplication, solving linear systems, Cholesky etc
- ▶ Sample from complicated distributions (such as truncated MVNs)

## Length of path required

- ▶ How long of a Markov chain do we need to approximate the posterior well?
- ▶ Informally, the higher the autocorrelations, the longer the path we will need
- ▶ Another performance metric: **effective sample size**, the equivalent number of **independent** samples (larger is better)

# Computational cost

- ▶ What is the computational cost? Two factors
  1. The cost of taking one step with  $\mathcal{P}$
  2. How long the Markov chain needs to be to make approximation “good”
- ▶ For the horseshoe, **both of these present challenges**
- ▶ Linear algebra with large matrices
- ▶ High autocorrelation

## Algorithmic developments

# Gibbs sampling for the horseshoe

State space  $\mathbf{X} = \mathbb{R}^p \times \mathbb{R}_+^p \times \mathbb{R}_+ \times \mathbb{R}_+$  with state-vector  $x = (\beta, \eta, \xi, \sigma^2)$ . Let  $D = \text{diag}(\eta_j^{-1})$

Typical computational approach: blocked Gibbs sampling (Polson et al (2012))

$$\eta \mid \beta, \sigma^2, \xi, z$$

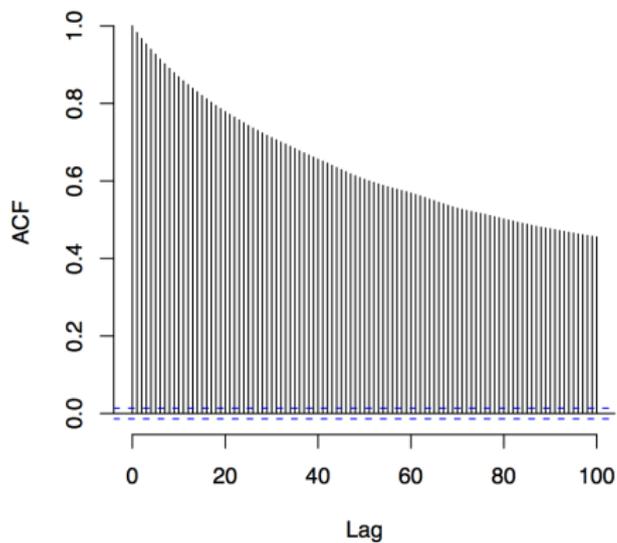
$$\xi \mid \beta, \sigma^2, \eta, z$$

$$(\beta, \sigma^2) \mid \eta, \xi, z$$

The algorithm is known to exhibit poor mixing for  $\xi$  (Polson et al. (2012))

# Mixing issues

Evidence of poor mixing for  $\xi$



## Remedy: Johndrow, Orenstein, B. (2018+)

- ▶ Our approach: more blocking

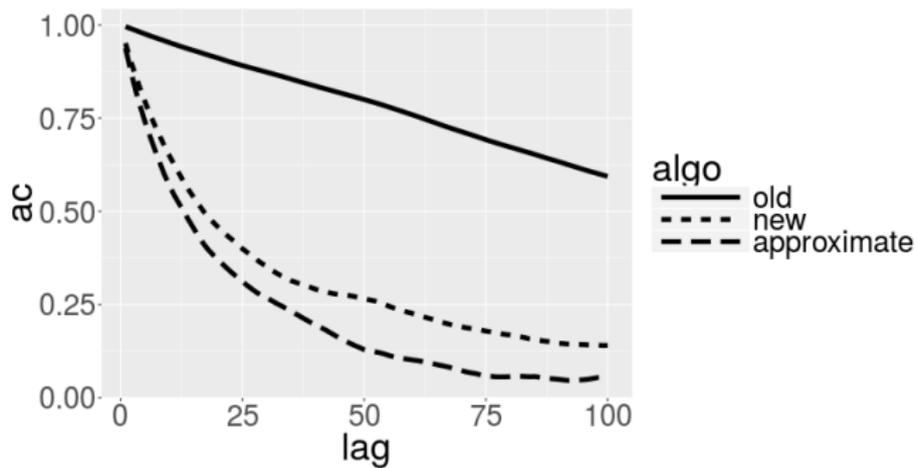
$$\begin{aligned}(\beta, \sigma^2, \xi) &| \eta, z \\ \eta &| (\beta, \sigma^2, \xi), z\end{aligned}$$

- ▶ The first step is done by sampling

$$\begin{aligned}\xi &| \eta, z \Rightarrow \text{Metropolis-within-Gibbs} \\ \sigma^2 &| \eta, \xi, z \Rightarrow \text{sample from Inverse-Gamma} \\ \beta &| \eta, \sigma^2, \xi, z \Rightarrow \text{sample from MVN}\end{aligned}$$

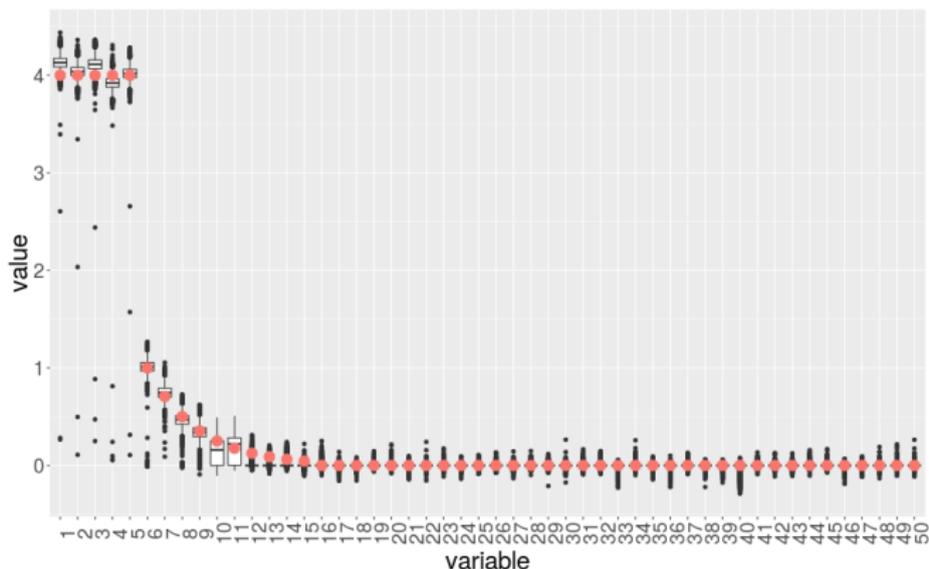
- ▶ The second step is done by sampling  $\eta_j$ s independently using an accurate rejection sampler

# Results: Autocorrelations for $\xi$

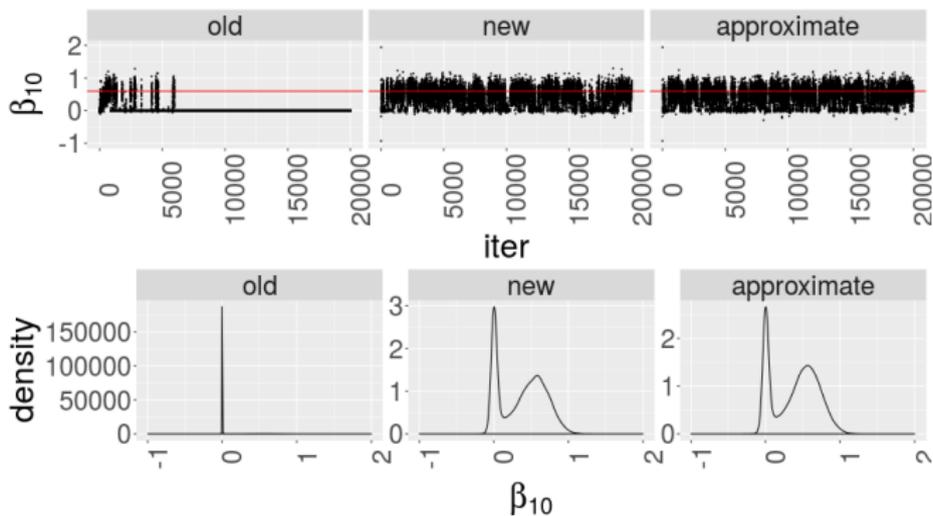


## Example

Simulation with  $N = 2,000$  and  $p = 20,000$ : first 50  $\beta_j$  (rest are zero). Posterior medians, 95 percent credible intervals, along with the truth.



## Results: Accuracy



The old algorithm often failed to identify components of  $\beta$  with bimodal marginals

This conveys uncertainty about whether  $\beta_j$  is a true signal, which is one of the nice features of taking a Bayesian approach to multiple testing

# Geometric ergodicity

**Theorem.** The blocked sampler above is **geometrically ergodic**.

Verify standard drift + minorization condition

1. **Foster–Lyapunov condition.** There exists a function  $V : \mathbf{X} \rightarrow [0, \infty)$  and constants  $0 < \gamma < 1$  and  $K > 0$  such that

$$(\mathcal{P}V)(x) \equiv \int V(y)\mathcal{P}(x, dy) \leq \gamma V(x) + K.$$

2. **Minorization.** For every  $R > 0$  there exists  $\alpha \in (0, 1)$  (depending on  $R$ ) such that, for  $\mathcal{S}(R) = \{x : V(x) < R\}$ ,

$$\sup_{x, y \in \mathcal{S}(R)} \|\mathcal{P}(x, \cdot) - \mathcal{P}(y, \cdot)\|_{TV} \leq 2(1 - \alpha).$$

## Geometric ergodicity

**Harris' Theorem** (Meyn & Tweedie; Hairer & Mattingley).

Let  $x = (\eta, \xi, \sigma^2, \beta)$  and  $\mathcal{P}$  the transition kernel. Also, let  $\mu$  be the invariant measure, i.e., the posterior.

Together, (1) and (2) imply,

$$\sup_{|\varphi| < 1+V} \int \varphi(y) (\mathcal{P}^n(x, y) - \mu(y)) dy \leq C \bar{\alpha}^n V(x),$$

for some  $\bar{\alpha} \in (0, 1)$ .

Geometric convergence in a weighted total variation norm

$(1 - \bar{\alpha})$  the spectral gap - larger implies faster convergence

## The exact algorithm

Blocking improves mixing, plus provably geometrically ergodic.

But what about the cost-per-step?

## Cost-per-iteration

Let's focus on the update of  $\beta$ :

$$\beta \mid \sigma^2, \xi, \eta, z \sim N \left( (W'W + (\xi^{-1}D)^{-1})^{-1} W'z, \sigma^2 (W^T W + (\xi^{-1}D)^{-1})^{-1} \right)$$

where  $D = \text{diag}(\eta_j^{-1})$ .

Usual Cholesky based sampler (Rue, 2001) for  $N(Q^{-1}b, Q^{-1})$  requires  $O(p^3)$  computation for non-sparse  $Q$ .

Highly prohibitive  $O(p^3)$  complexity per iteration when  $p \gg N$ .

## (Partial) Remedy

In B., Chakraborty, Mallick (2016), we propose an alternative exact sampler with  $O(N^2 p)$  complexity.

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(i) Sample  $u \sim N(0, \xi^{-1}D)$  and  $f \sim N(0, I_N)$  indep.

(ii) Set  $v = Wu + f$

(iii) Solve  $M_\xi v^* = (z/\sigma - v)$  where  $M_\xi = I_N + \xi^{-1}WDW'$

(iv) Set  $\beta = \sigma(u + \xi^{-1}DW'v^*)$

---

(iii) is the costliest step taking  $\max\{O(N^2 p), O(N^3)\}$  steps. Significant savings when  $p \gg N$ .

## Cost-per-iteration

However, still  $O(N^2p)$  computation.  $N$  can be in the order of tens of thousands in GWAS studies.

The remaining bottleneck is only in calculating

$$M_\xi = I_N + \xi^{-1}WDW'$$

which is needed by the updates for  $\beta$ ,  $\sigma^2$ , and  $\xi$

Our proposal: replace  $WDW'$  with a cheaper and accurate approximation

## Approximations in MCMC

# Approximation

- ▶ Horseshoe is designed to shrink most coordinates of  $\beta$  toward zero... So many of the  $(\xi\eta_j)^{-1}$  will typically be tiny at any iteration
- ▶ Choose a “small” threshold  $\delta$ , approximate  $M_\xi$  by

$$M_{\xi,\delta} = I_N + \xi^{-1} W_S D_S W_S', \quad S = \{j : \xi^{-1} \eta_j^{-1} > \delta\}$$

where  $W_S$  is the sub-matrix consisting of columns in the set  $S$ , etc

- ▶ Carefully replace all calculations involving  $M_\xi$  with  $M_{\xi,\delta}$
- ▶ Reduces cost per step to  $Ns^2 \vee Np$ , where  $s = |S|$

**Note:** this is different from setting some  $\beta_j = 0$  at each scan.  $\beta$  is still being drawn from a non-singular MVN.

# Perturbations in MCMC

- ▶ A general strategy to reduce cost-per-step is to replace the exact transition kernel  $\mathcal{P}$  with an “approximation”  $\mathcal{P}_\epsilon$
- ▶ Some other examples - replace a non-standard density with its best approximation from a standard family, divide-conquer...
- ▶  $\mathcal{P}_\epsilon$  still a Markov chain
- ▶ Question: what can we say about finite-time averages from the approximate chain? In other words, is

$$\mu\varphi \approx n^{-1} \sum_{k=0}^{n-1} \varphi(X_k^\epsilon)$$

for  $X_k^\epsilon \sim \nu \mathcal{P}_\epsilon^k$ ?

## Literature review

- ▶ Early reference on perturbation bounds: Mitrophanov (2005), for uniformly ergodic chains
- ▶ Renewed interest in recent years (Alquier et al. 2014, Pillai & Smith (2015), Rudolf & Schweizer (2018), Johndrow & Mattingley (2018)) - extensions to unbounded state-spaces
- ▶ Most applications pertain to “tall data”, i.e., lots of samples (Bardenet, Doucet, Holmes (2017))
- ▶ Ours is one of the first applications for large  $N$  and  $p$  with potentially  $p \gg N$

## A new general perturbation bound

We show that

$$\mathbf{E} \left( \frac{1}{n} \sum_{k=0}^{n-1} \varphi(X_k^\epsilon) - \mu\varphi \right)^2$$

can be “controlled” (skipping exact bounds) if

1. There exists  $K_\epsilon > 0$  and  $\gamma_\epsilon \in (0, 1)$  such that

$$(\mathcal{P}_\epsilon V)(x) \leq \gamma_\epsilon V(x) + K_\epsilon,$$

that is  $V$  is also Lyapunov for  $\mathcal{P}_\epsilon$ .

2. The approximate kernel  $\mathcal{P}_\epsilon$  satisfies

$$\sup_{x \in \mathbf{X}} \|\mathcal{P}(x, \cdot) - \mathcal{P}_\epsilon(x, \cdot)\|_{TV} \leq \frac{\epsilon}{2}.$$

## Application to Horseshoe sampler

Recall our approximation step replaces  $M_\xi = I_N + \xi^{-1}WDW'$  with  $M_{\xi,\delta} = I_N + \xi^{-1}WD_\delta W'$ .

We show that this approximation achieves

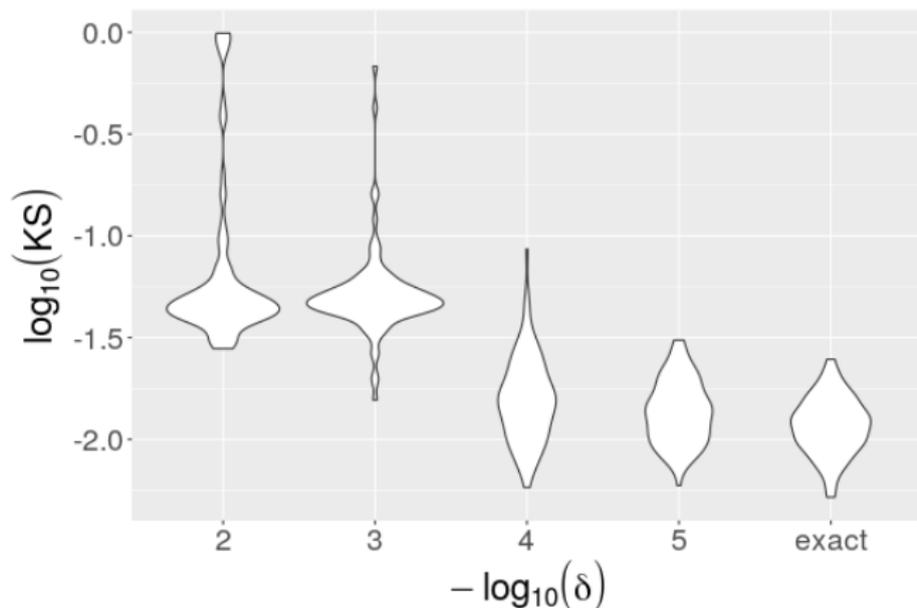
$$\sup_x \|\mathcal{P}(x, \cdot) - \mathcal{P}_\delta(x, \cdot)\|_{TV}^2 \leq \delta \|W\|^2 [4N(\|z\|^2/b_0) + 9] + \mathcal{O}(\delta^2)$$

for any small fixed threshold  $\delta$ .

Satisfies conditions of our general theorem.

## Application to Horseshoe sampler

Practically: we recommend  $\delta = 10^{-4}$  or  $10^{-5}$  and have observed no advantages from smaller values.



**Figure:** Average KS distance between the marginals of 100 entries of  $\beta$  from the exact and approximate algorithm for  $N = 1000$  and  $p = 10000$

## Varying threshold

- ▶ Using a fixed threshold  $\epsilon$  results in an asymptotic bias proportional to  $\sqrt{\epsilon}/(1 - \bar{\alpha})$ , where recall  $\bar{\alpha}$  quantifies rate of convergence of the exact chain
- ▶ More room to use approximations when the exact chain mixes rapidly, i.e.,  $\sqrt{\epsilon}$  is small compared to the spectral gap  $(1 - \bar{\alpha})$  of the exact chain
- ▶ The asymptotic bias can be eliminated by using a decreasing schedule of approximation parameters  $(\epsilon_k)$  - need to satisfy  $\epsilon_k \rightarrow 0$  “sufficiently fast” (summability condition)
- ▶ Reminiscent of conditions for stochastic gradient or Langevin dynamics

# Simulation studies

The results that follow use a common simulation structure

$$\begin{aligned}w_i &\stackrel{iid}{\sim} N_p(0, \Sigma) \\z_i &\sim N(w_i\beta, 4) \\ \beta_j &= \begin{cases} 2^{-(j/4-9/4)} & j < 24 \\ 0 & j > 23 \end{cases},\end{aligned}$$

So there are always “small” and “large” signals, and true nulls  
We consider both  $\Sigma = I$  (**independent design**) and  $\Sigma_{ij} = 0.9^{|i-j|}$   
(**correlated design**)

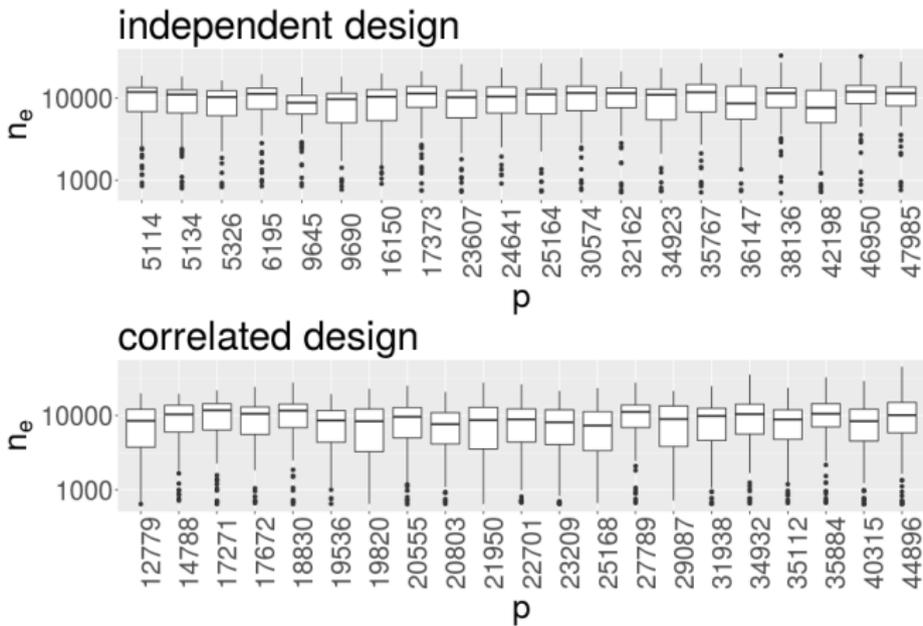
## Effective sample size

Recall **effective sample size**  $n_e$ , a measure of the number of **independent** samples your Markov path is “worth”

If  $n_e = n$  then your MCMC is giving essentially independent samples (like vanilla Monte Carlo)

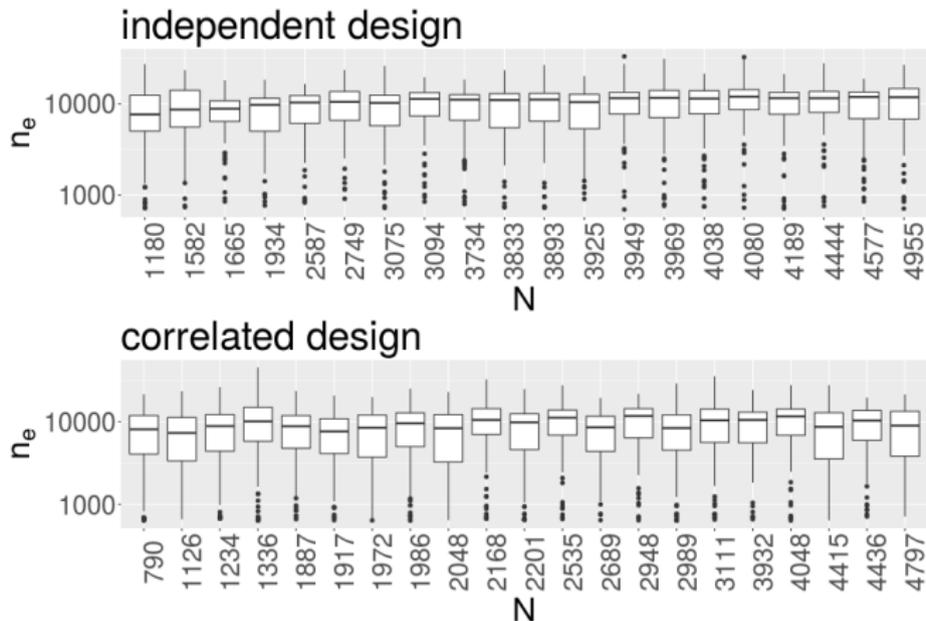
If  $n_e \ll n$  then your MCMC has very high autocorrelations, need very long path to get good approximation to posterior

# Mixing as $p$ increases



Effective sample sizes are essentially independent of  $p$ , even when the design matrix is highly correlated

# Mixing as $N$ increases



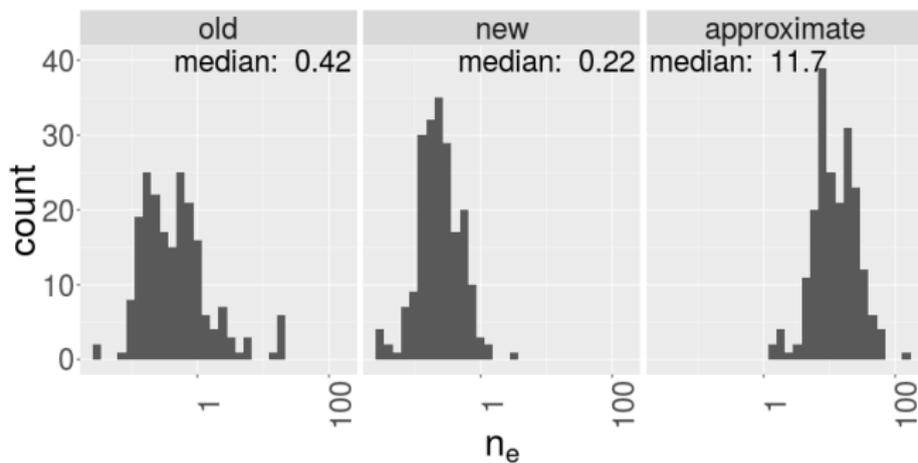
Effective sample sizes are essentially independent of  $N$ , even when the design matrix is highly correlated

## Effective samples per second

Recall **effective sample size**  $n_e$ , a measure of the number of **independent** samples your Markov path is “worth”

So if  $t$  is computation time in seconds, **effective samples per second**  $n_e/t$  is an empirical measurement of overall computational efficiency

## Results: Effective samples per second



The approximate algorithm is **fifty times** more efficient when  $N = 2,000$  and  $p = 20,000$

# Conclusion

Computational cost for MCMC shouldn't massively differ from alternatives designed for the same problem

But making the algorithm fast takes work, often problem-specific

More thrust on “computing” posteriors that we know have “nice” properties

Approximations in MCMC seem a promising direction to speed-up computation

A step towards rigorous quantification of approximation error

## References

- ▶ Bhattacharya, A., Chakraborty, A., Mallick, B. (2016). Fast sampling with Gaussian scale-mixture priors in high-dimensional regression. *Biometrika* arXiv preprint arXiv:1506.04778
- ▶ Johndrow, J. E., Orenstein, P., & Bhattacharya, A. (2018). Bayes Shrinkage at GWAS scale: Convergence and Approximation Theory of a Scalable MCMC Algorithm for the Horseshoe Prior arXiv preprint arXiv:1705.00841.
- ▶ Pallavi Ray, Anirban Bhattacharya, and Debdeep Pati. <https://arxiv.org/abs/1902.04701>

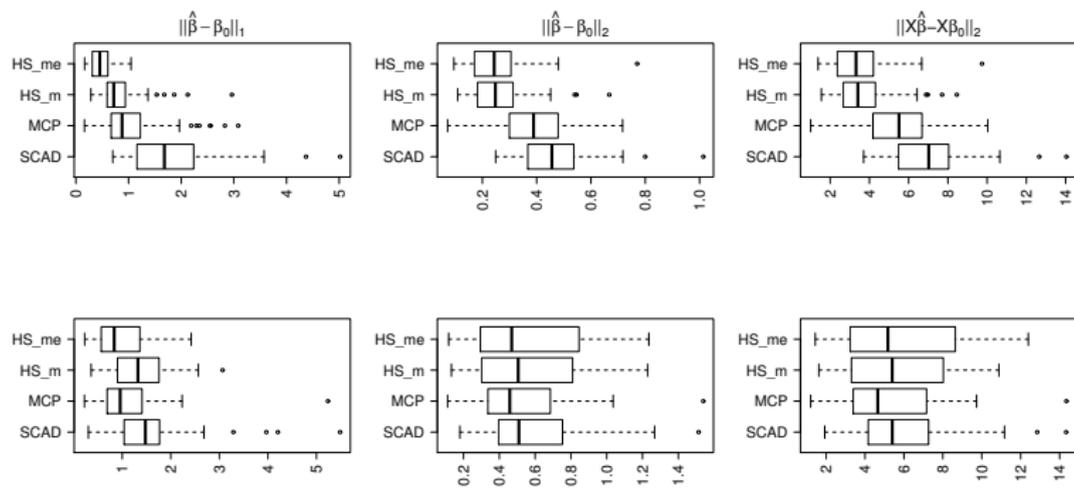
Thank You

Performance in  $p \gg n$  settings

# Data generation

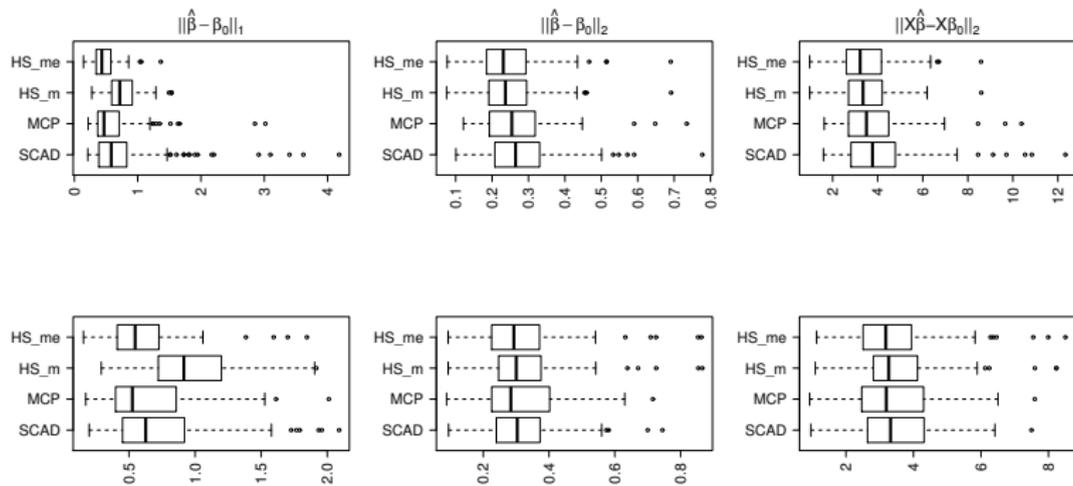
- ▶ Replicated simulation study with horseshoe prior (Carvalho et al. (2010))
- ▶  $n = 200$  &  $p = 5000$ . True  $\beta_0$  has 5 non-zero entries and  $\sigma = 1.5$
- ▶ Two signal strengths:
  - (i) *weak* -  $\beta_{0S} = \pm(0.75, 1, 1.25, 1.5, 1.75)^T$
  - (ii) *moderate* -  $\beta_{0S} = \pm(1.5, 1.75, 2, 2.25, 2.5)^T$
- ▶ Two types of design matrix:
  - (i) *Independent* -  $X_j$  i.i.d.  $N(0, I_p)$
  - (ii) *compound symmetry* -  $X_j$  i.i.d.  $N(0, \Sigma)$ ,  $\Sigma_{jj'} = 0.5 + 0.5\delta_{jj'}$
- ▶ Summary over 100 datasets

# Weak signal case



Estimation performance: Boxplots of  $\ell_1$ ,  $\ell_2$  and prediction error across 100 simulation replicates.  $HS_{me}$  and  $HS_m$  are posterior point wise median and mean for the horeshoe prior. Top row: Independent covariates, Bottom row: Compound symmetry

# Moderate signal case



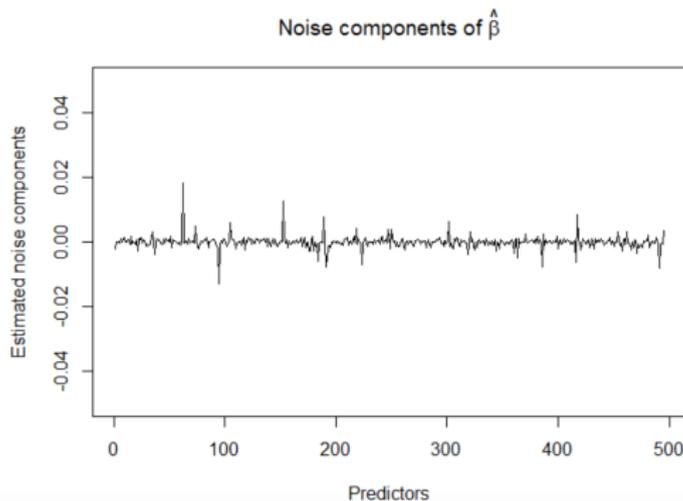
Estimation performance: Boxplots of  $\ell_1, \ell_2$  and prediction error across 100 simulation replicates.  $HS_{me}$  and  $HS_m$  are posterior point wise median and mean for the horeshoe prior. Top row: Independent covariates, Bottom row: Compound symmetry

# Frequentist coverage of 95% credible intervals

p	500								
Design	Independent			Comp Symm			Toeplitz		
	HS	LASSO	SS	HS	LASSO	SS	HS	LASSO	SS
Signal Coverage	93 <sub>1.0</sub>	75 <sub>12.0</sub>	82 <sub>3.7</sub>	95 <sub>0.9</sub>	73 <sub>4.0</sub>	80 <sub>4.0</sub>	94 <sub>4.0</sub>	80 <sub>7.0</sub>	79 <sub>5.6</sub>
Signal Length	42	46	41	85	71	75	86	79	74
Noise Coverage	100 <sub>0.0</sub>	99 <sub>0.8</sub>	99 <sub>1.0</sub>	100 <sub>0.0</sub>	98 <sub>1.0</sub>	99 <sub>0.8</sub>	98 <sub>1</sub>	98 <sub>1.0</sub>	99 <sub>0.6</sub>
Noise Length	2	43	40	4	69	73	5	78	73

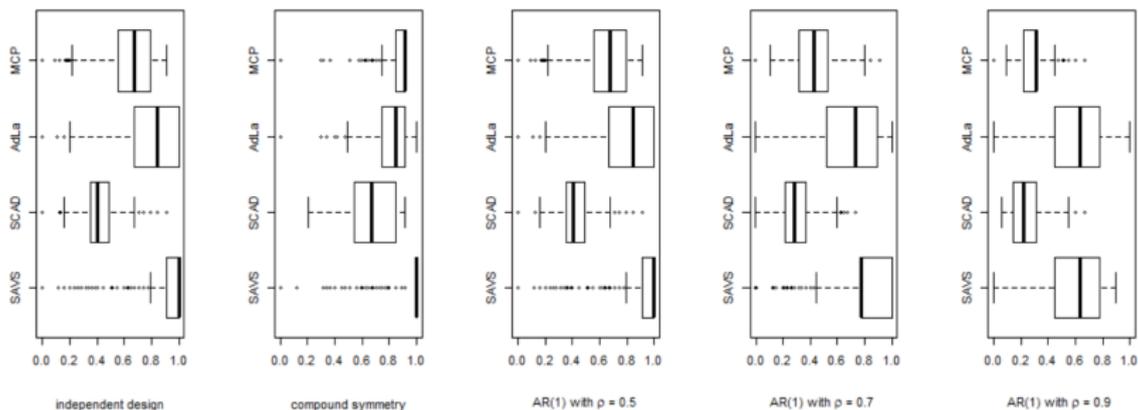
Frequentist coverages (%) and  $100 \times$  lengths of point wise 95% intervals. Average coverages and lengths are reported after averaging across all signal variables (rows 1 and 2) and noise variables (rows 3 and 4). Subscripts denote  $100 \times$  standard errors for coverages. LASSO and SS respectively stand for the methods in van de Geer et al. (2014) and Javanmard & Montanari (2014). The intervals for the horseshoe (HS) are the symmetric posterior credible intervals.

# Variable selection by postprocessing



$$Q(\beta) = \frac{1}{2} \|\mathbf{X}\hat{\beta} - \mathbf{X}\beta\|_2^2 + \sum_{j=1}^p \mu_j |\beta_j|, \quad \mu_j = |\hat{\beta}_j|^{-2}.$$

# Variable selection performance



SAVS: Variable selection by post-processing the posterior mean from the HS prior. Plot of Mathew's correlation coefficient (MCC) over 1000 simulations for various methods. MCC values closer to 1 indicate better variable selection performance.