

Abstract

Bayesian statistical models are attractive in the Big Data setting, but Markov Chain Monte Carlo methods – the cornerstone of modern Bayesian computation – do not extend easily to the cluster setting, where the data is too big to fit on one machine. We present a novel scheme to parallelize MCMC with no synchronization or locking, avoiding typical performance bottlenecks, particularly in settings where the number of model parameters grows with data size.

Big Data: the problem

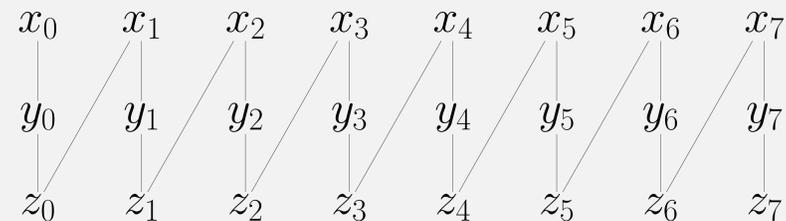
Data: *too big to fit on one machine*

- Lives on a Hadoop cluster: fault-tolerant distributed storage
 - Big Data \iff Big Hardware
- Computational cost of *1 data point* \approx cost of *all N points*
 - Data *must* be operated on *in parallel*
 - Need parallel algorithms that scale *superlinearly*
- Model dimensionality often grows with N

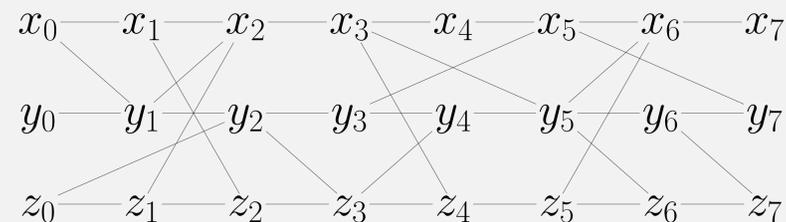
Previous Work on Scalable Bayesian Inference

- Data that fits on *one machine*
 - Approximate Methods: Variational Bayes [2], ABC [6]
 - Exact Methods: Langevin Diffusion based methods [4]
- Data *too big for one machine*
 - *Consensus Monte Carlo* [5]: for models where dimensionality doesn't grow with N (exact for some models)
 - *Hogwild*: [3] asynchronous stochastic gradient descent, point estimation only, widely used in machine learning

Standard Gibbs Sampling



Asynchronous Gibbs Sampling



The Algorithm

1. Start with a *latent variable* model, partition data among workers
2. Run Gibbs steps on each worker
 - Condition on *most recent local variables*
 - Condition on *most recent known non-local* variables, which *may be out of date* in the cluster as a whole
 - *Draw* samples, *transmit* to other workers
 - Never stop, never synchronize, never wait

Implementation

- Highly non-trivial: not expressible in the MapReduce paradigm (typical high-level parallel computation framework used in Hadoop)
- Expressible in shared memory and actor models of parallelism
- Written in Scala, a modern language well-suited to parallel use cases
- Code available via eBay Software Foundation on GitHub

Convergence

Two kinds of algorithms: different communication between workers

- *Exact Asynchronous Gibbs*: apply Metropolis-Hastings correction
- *Approximate Asynchronous Gibbs*: accept all updates

Theorem: *Exact Asynchronous Gibbs converges*

- Intuitive sketch of proof
 1. *Instant communication \implies exact algorithm converges*
 - Define a Markov chain that (1) selects a worker, (2) selects a full conditional, (3) proposes a new state from that full conditional at every worker, (4) performs a Metropolis step on each worker
 - Proof via detailed balance, workers accept/reject independently
 2. *Exact algorithm converges \implies asynchronous convergence*
 - Proof via result on convergence of asynchronous algorithms

Approximate Asynchronous Gibbs: Noisy Monte Carlo [1]

- Replace Metropolis-Hastings ratio with *biased estimator* – 1
- Perform *diagnostic check* at runtime to ensure *bias is small*

Future Work

- Better understanding of Noisy Monte Carlo [1] approximations
- Implement algorithm in *MPI* for use on traditional supercomputers

Results

Gaussian Process Regression: *71,500 latent variables* (toy problem)
– *Correct answer* in *20 minutes* on 143-core cluster

Large hierarchical model: *1,000,000 latent variables* (real-world)
– *Same answer* as sequential-scan Gibbs (up to Monte Carlo error)

Standard Gibbs Sampler	12 hrs
8-core machine, high memory, problem-specific parallelization	
Asynchronous Gibbs	1 hr
160-core cluster	

References

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- [6] S. Tavaré, D. J. Balding, R. C. Griffiths, and P. Donnelly. Inferring coalescence times from DNA sequence data. *Genetics*, 145(2):505–518, 1997.

Acknowledgments

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