

# Project Proposal

Pramoda Jayasinghe

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The aim of this project is to evaluate the performance of different Markov Chain Monte–Carlo (MCMC) methods for computing posterior distributions. Achieving this can be expressed as two sub objectives:

1. Implementing the code for two selected algorithms using R.
2. Evaluating the performance of these algorithms using effective sample size (ESS) and exploration quality (EQ) (Ballnus et al., 2017; Turner and Neal, 2017).

## 1 Implementation

Two MCMC algorithms are selected considering the time available for this project. The commonly known Metropolis–Hastings (MH) and more advanced Parallel Tempering (PT) algorithms are used in the comparison of this project. These methods are implemented for 2 cases.

1. A simple Bernoulli distribution with a continuous Uniform parameter, and
2. a finite mixture problem.

## 2 Performance Evaluation

To evaluate the performance of the algorithms, can be evaluated separately for the two problems. To this end, ESS and EQ methods can be used. Using two different tools to benchmark will allow for a comparison between the benchmarking tools too. The computation time can also be measured as an optional benchmarking tool.

## References

- Ballnus, B., S. Hug, K. Hatz, L. Görlitz, J. Hasenauer, and F. J. Theis (2017, dec). Comprehensive benchmarking of Markov chain Monte Carlo methods for dynamical systems. *BMC Systems Biology* 11(1), 63.
- Turner, R. and B. Neal (2017, dec). How well does your sampler really work?