

# OxWaSP Module 1: Adaptive MCMC

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## 1 Introduction to Adaptive MCMC - the AM algorithm

MCMC algorithms allow sampling from complicated, high-dimensional distributions. Choice of the proposal distribution (from which samples are taken in an attempt to approximate sampling from the target distribution  $\pi$ ) determines the ability of the algorithm to explore the parameter space fully and hence draw a good sample. Adaptive MCMC algorithms tackle this challenge by using samples already generated to learn about the target distribution; they push this knowledge back to the choice of proposal distribution iteratively.

This project explores adaptive MCMC algorithms existing in the literature that use covariance estimators to improve convergence to a target distribution supported on a subset of  $\mathbb{R}^d$ . In this schema we learn about the target distribution  $\pi$  through estimation of its correlation structure from the MCMC samples. We use this correlation structure to improve our estimate of the target.

We first implement an adaptive MCMC algorithm AM (Haario et al., 2001) which is a modification of the random walk Metropolis-Hastings algorithm. In AM the proposal distribution is updated at time  $t$  to be a normal distribution centered on the current point  $X_{t-1}$  with covariance  $C_t(X_0, \dots, X_{t-1})$  that depends on the whole history of the chain. The use of historic states means the resulting chain is non-markovian, and reversibility conditions are not satisfied. Haario et al show that, with a small update to the usual Metropolis-Hastings acceptance probability, the right ergodic properties and correct simulation of the target distribution none the less remain. The probability with which to accept candidate points in the chain becomes:

$$\alpha(X_{t-1}, Y) = \min \left( 1, \frac{\pi(Y)}{\pi(X_{t-1})} \right)$$

With  $C_t$  given by:

$$C_t = s_d \text{cov}(X_0, \dots, X_{t-1}) + s_d \epsilon I_d$$

Here  $\text{cov}()$  is the usual empirical covariance matrix, and the parameter  $s_d = \frac{2.4^2}{d}$  (Gelman et al., 1996).  $\epsilon$  is chosen to be very small compared to

the subset of  $\mathbb{R}^d$  upon which the target function is supported. The AM algorithm is computationally feasible due to recursive updating of the covariance matrix on acquisition of each new sample through the relation:

$$C_{t+1} = \frac{t-1}{t}C_t + \frac{s_d}{t}(t\bar{X}_{t-1}\bar{X}_{t-1}^T - (t+1)\bar{X}_t\bar{X}_t^T + X_tX_t^T + \epsilon I_d)$$

with the mean calculated recursively by:

$$\bar{X}_{t+1} = \frac{t\bar{X}_t + X_{t+1}}{t+1}$$

Because of the instability of the covariance matrix, to implement the adaptivity we first run the algorithm with no change to the covariance of the proposal distribution. The adaptation starts at a user defined point in time, and until this time the covariance of the proposal is chosen to represent our best knowledge of the target distribution.

## 2 An example - testing the AM algorithm

We now numerically test the AM algorithm. We have used two different target distributions: a correlated Gaussian distribution  $N(0, \Sigma)$  and a "banana"-shaped distribution ((Roberts and Rosenthal, 2009)) given by:

$$f_B(x_1, \dots, x_d) \propto \exp \left[ -x_1^2/200 - \frac{1}{2} (x_2 + Bx_1^2 - 100B)^2 - \frac{1}{2} (x_3^2 + x_4^2 + \dots + x_d^2) \right]$$

$B > 0$  is the "bananicty" constant (set to 0.1 throughout) and  $d$  is the dimension. We have chosen the correlated Gaussian distribution as targetting this demonstrates how the use of empirical covariance improves convergence - we learn the target's covariance as we move through steps of the MCMC. The banana-shaped distribution is an additional example with an irregular shape. We use this to test the ability of the markov chain to fully explore the state space with and without adaption. We first run our implementation of the AM algorithm targetting  $N(0, \Sigma)$  with

```
d = 8
n_iter = 200      # Total number of iterations
x_1 = rep(5,d)    # Vector of initial values
t_adapt = 50      # When to start adapting
adapt = "None"    # Choose the type of adaptation.

X_MH = mcmc(target = pi_norm_corr,
             n_iter = n_iter,
             x_1 = x_1,
             adapt=adapt)
```

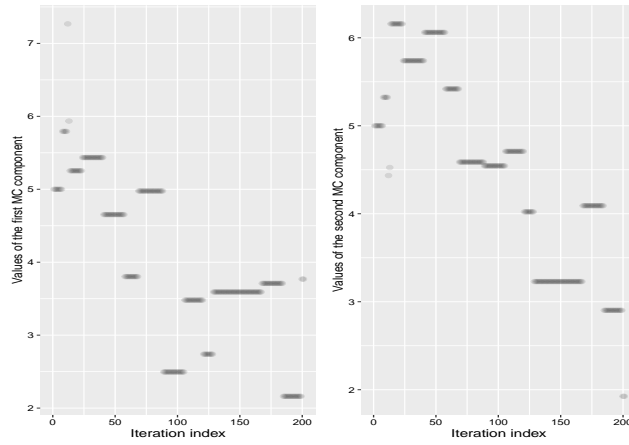


Figure 1: The first (left) and second(right) components of the markov chain resulting from Metropolis-Hastings (MH) on a correlated 8-dimensional Gaussian.

```
## Running MCMC targeting pi_norm_corr with 200 iterations.
## Adaptation algorithm is: None .
## Covariance matrix estimator is: Sample covariance .
## MCMC finished in 0.1043711 seconds.
## The last acceptance rate was: 0.075
```

```
adapt = "AM"           # Choose the type of adaptation.
cov_estimator="Sample covariance" # Choose the type of covariance matrix estimator.

X_AM = mcmc(target = pi_norm_corr,
            n_iter = n_iter,
            x_1 = x_1,
            adapt=adapt,
            t_adapt = t_adapt,
            cov_estimator=cov_estimator
            )

## Running MCMC targeting pi_norm_corr with 200 iterations.
## Adaptation algorithm is: AM .
## Covariance matrix estimator is: Sample covariance .
## MCMC finished in 1.92934 seconds.
## The last acceptance rate was: 0.19

plot6=plotIterations(
    X_AM$X[,1],
    n_iter,
```

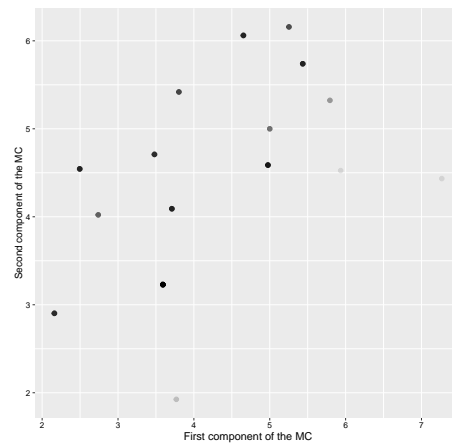


Figure 2: This is the second graph

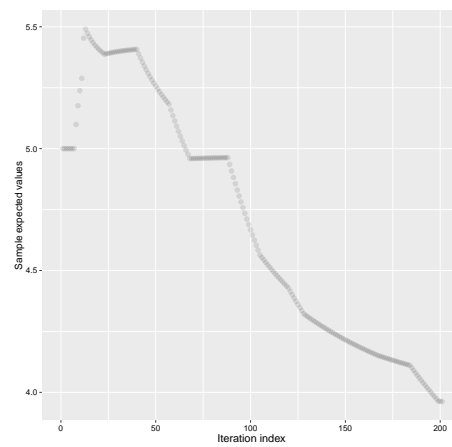


Figure 3: This is the third graph

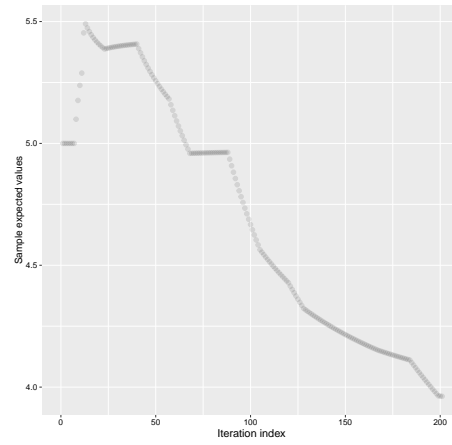
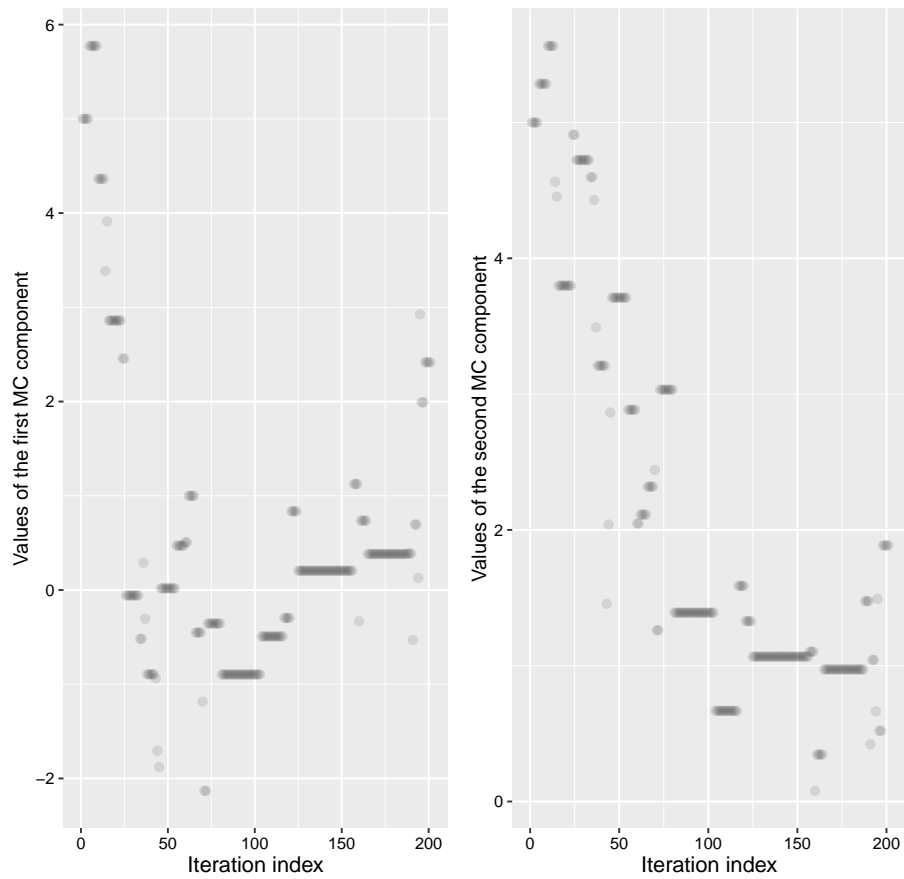


Figure 4: This is the fourth graph

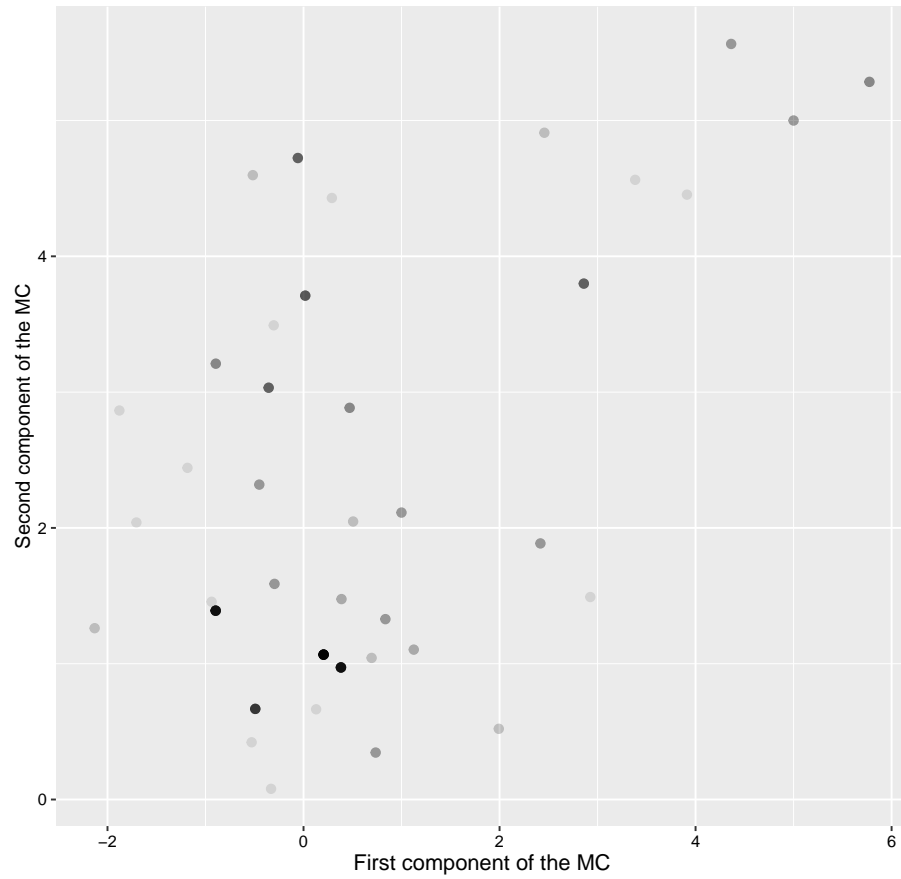
```

        title="Values of the first MC component"
    )
plot7=plotIterations(
    X_AM$X[,2],
    n_iter,
    title="Values of the second MC component"
)
grid.arrange(plot6, plot7, ncol=2)

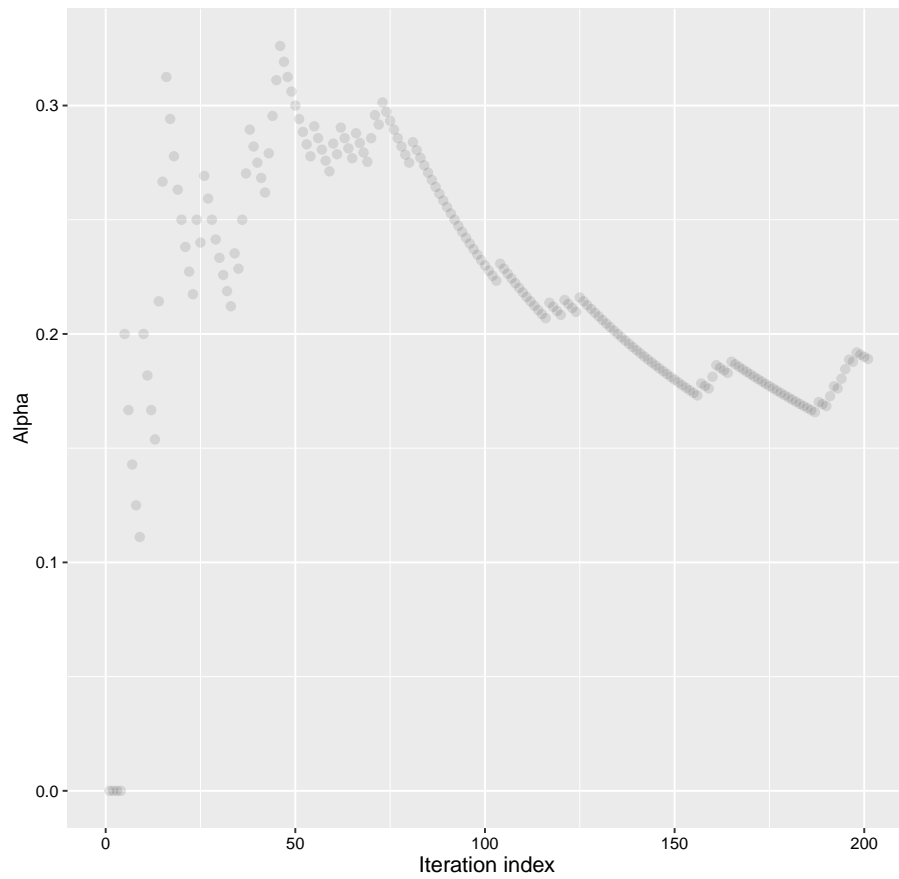
```



```
plot8=plotComponents(
    X_AM$X[,1],
    X_AM$X[,2],
    Xtitle = "First component of the MC",
    Ytitle = "Second component of the MC"
)
plot8
```

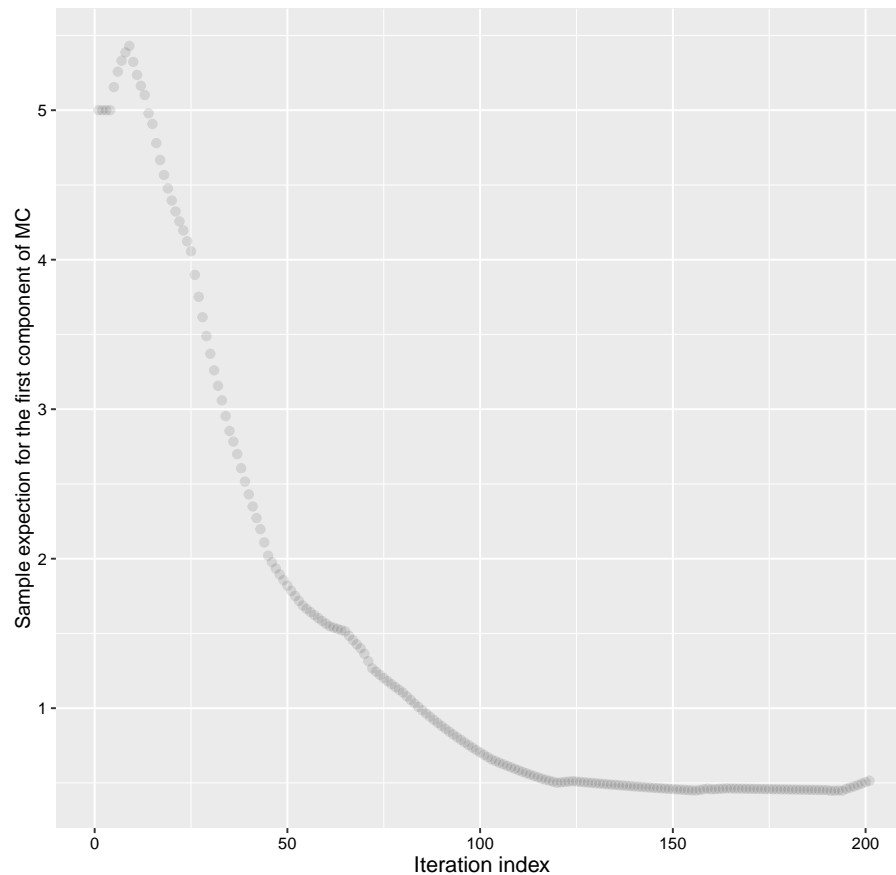


```
plot9=plotIterations(X_AM$acceptance_rates, n_iter, "Alpha")  
plot9
```



```
plot10=plotIterations(X_AM$sample_mean[,1], n_iter, "Sample expectation for the first component")
plot10
```





AM is able to explore the state space, adapt properly and settle down to rapid mixing. We see an improved acceptance rate.

```
x_1 = rep(5,2)      # Vector of initial values
target = pi_banana

X_MH_banana = mcmc(target = target,
  n_iter = n_iter,
  x_1 = x_1,
  adapt="None"
)

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: None .
## Covariance matrix estimator is: Sample covariance .
## MCMC finished in 0.04179502 seconds.
```

```

## The last acceptance rate was: 0.24

plot11=plotComponents(
    X_MH_banana$X[,2],
    X_MH_banana$X[,1],
    Xtitle = "First component of the MC",
    Ytitle = "Second component of the MC"
)

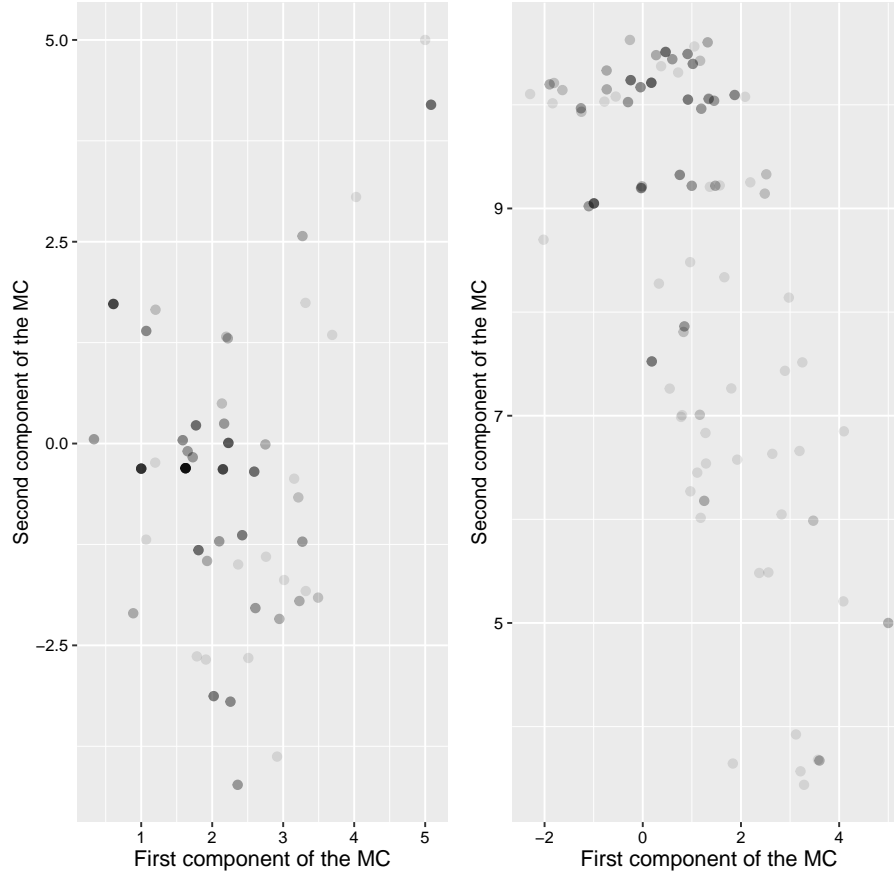
X_AM_banana = mcmc(target = target,
    n_iter = n_iter,
    x_1 = x_1,
    t_adapt = t_adapt,
    adapt="AM"
)

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: AM .
## Covariance matrix estimator is: Sample covariance .
## MCMC finished in 1.709115 seconds.
## The last acceptance rate was: 0.41

plot12=plotComponents(
    X_AM_banana$X[,2],
    X_AM_banana$X[,1],
    Xtitle = "First component of the MC",
    Ytitle = "Second component of the MC"
)

grid.arrange(plot11, plot12, ncol=2)

```



We now explore a slight modification to the adaptation scheme AM2(Roberts and Rosenthal, 2009), that uses stochastic stabilisation rather than the numerical stabilisation of AM. Roberts and Rosenthal use a mixture of Gaussians as the proposal distribution: with proportion  $\beta$  a normal uncorrelated distribution is mixed with a correlated normal distribution.

$$Q_n(x, \cdot) = (1 - \beta)N(x, s_d \Sigma_n) + \beta(N(x, (0.1^2)I_d/d)$$

TODO Discuss the problems with starting dimension here.

```
x_1 = rep(5,8)      # Vector of initial values
target = pi_banana8

X_AM2_banana = mcmc(target = target,
  n_iter = n_iter,
  x_1 = x_1,
```

```

        adapt="AM2"
    )

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: AM2 .
## Covariance matrix estimator is: Sample covariance .
## MCMC finished in 0.03528547 seconds.
## The last acceptance rate was: 0.28

plot13=plotIterations(
    X_AM_banana$X[,1],
    n_iter,
    title="Values of the first MC component"
)
plot14=plotIterations(
    X_AM2_banana$X[,1],
    n_iter,
    title="Values of the first MC component"
)

grid.arrange(plot13, plot14, ncol=2)

```

```

x_1 = rep(5,8)      # Vector of initial values
cov_estimator1="Shrinkage estimator"
cov_estimator2="Thresholding estimator"
target = pi_banana8

X_MH_banana = mcmc(target = target,
    n_iter = n_iter,
    x_1 = x_1,
    adapt="None"
)

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: None .
## Covariance matrix estimator is: Sample covariance .
## MCMC finished in 0.06426764 seconds.
## The last acceptance rate was: 0.065

X_AM_banana = mcmc(target = target,
    n_iter = n_iter,
    x_1 = x_1,
    t_adapt = t_adapt,
    adapt="AM"
)

```

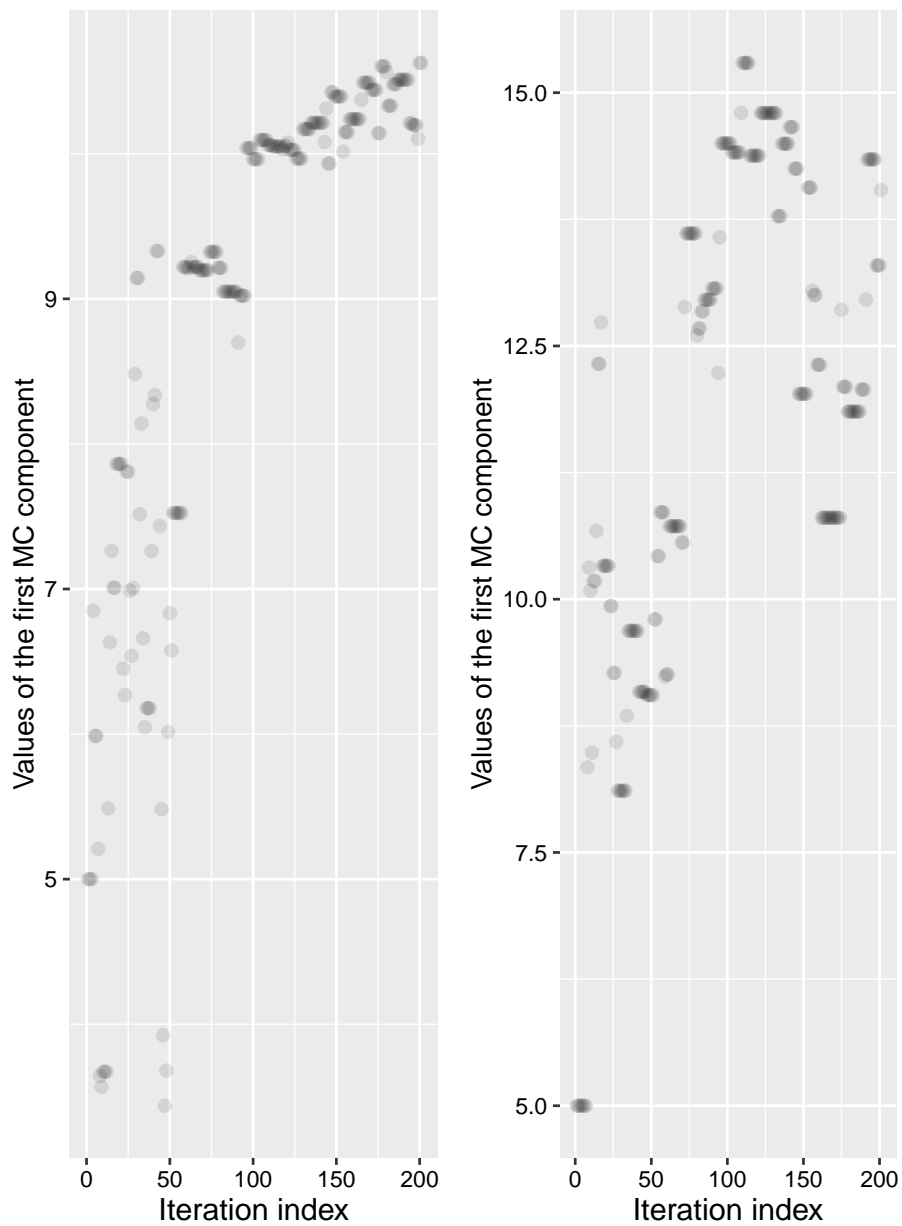


Figure 5: The first component of the 8-dimensional banana shaped distribution targetted using AM (left) and AM2 (right).

```

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: AM .
## Covariance matrix estimator is: Sample covariance .
## MCMC finished in 1.799823 seconds.
## The last acceptance rate was: 0.175

X_AM_banana_sh= mcmc(target = target,
                     n_iter = n_iter,
                     x_1 = x_1,
                     adapt="AM",
                     t_adapt = t_adapt,
                     cov_estimator=cov_estimator1
                     )

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: AM .
## Covariance matrix estimator is: Shrinkage estimator .Estimating optimal shrinkage intens
##
##
## MCMC finished in 2.159014 seconds.
## The last acceptance rate was: 0.24

X_AM2_banana_sh = mcmc(target = target,
                       n_iter = n_iter,
                       x_1 = x_1,
                       adapt="AM2",
                       cov_estimator=cov_estimator1
                       )

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: AM2 .
## Covariance matrix estimator is: Shrinkage estimator .
## MCMC finished in 0.02885199 seconds.
## The last acceptance rate was: 0.27

X_AM_banana_th= mcmc(target = target,
                     n_iter = n_iter,
                     x_1 = x_1,
                     adapt="AM",
                     t_adapt = t_adapt,
                     cov_estimator=cov_estimator2
                     )

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: AM .
## Covariance matrix estimator is: Thresholding estimator .
## MCMC finished in 0.2311356 seconds.
## The last acceptance rate was: 0.235

```

```

X_AM2_banana_th = mcmc(target = target,
  n_iter = n_iter,
  x_1 = x_1,
  adapt="AM2",
  cov_estimator=cov_estimator2
)

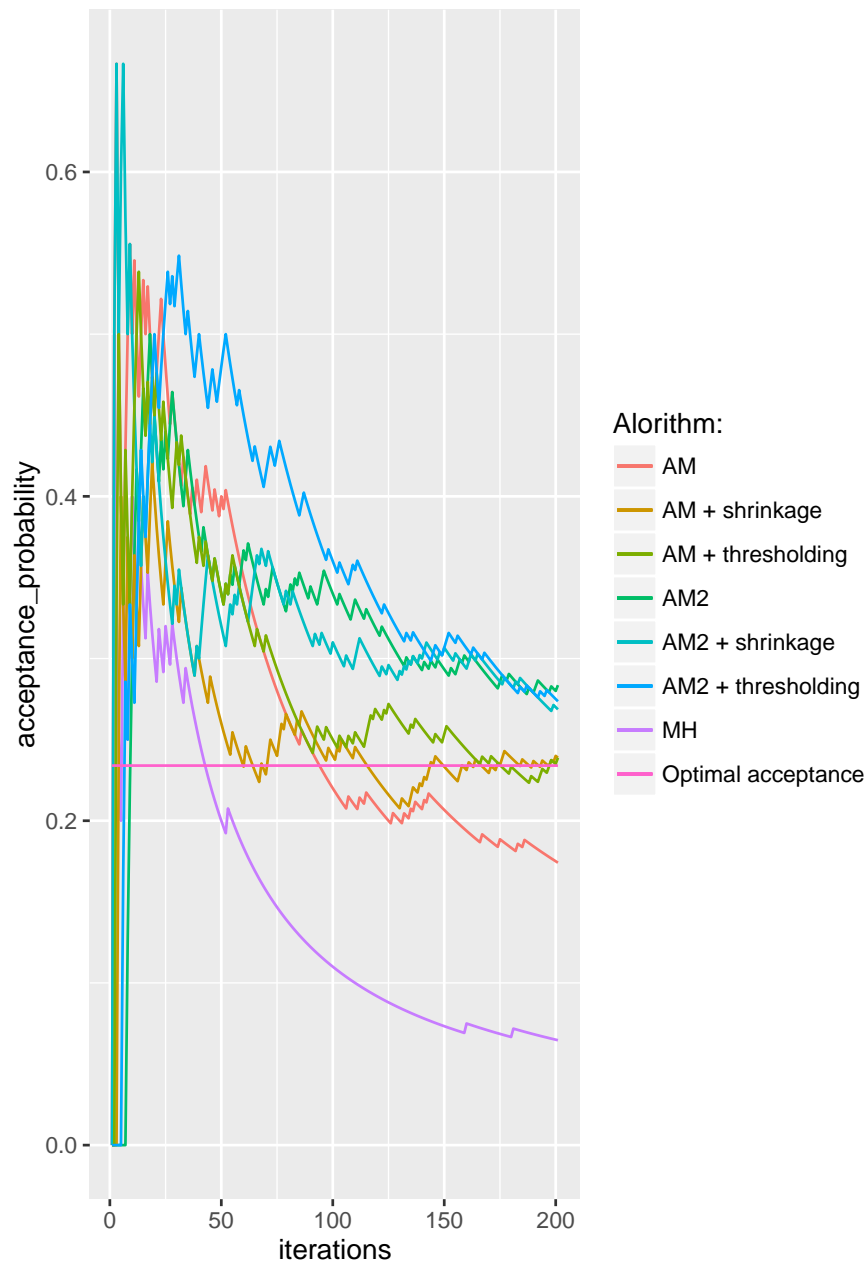
## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: AM2 .
## Covariance matrix estimator is: Thresholding estimator .
## MCMC finished in 0.03438878 seconds.
## The last acceptance rate was: 0.275

iteration=seq(from= 1, to=n_iter+1, by=1)
#data=data.frame(iteration_count=iteration, am_alpha=X_AM_banana_sh$acceptance_rates)

df1<-data.frame(iterations=iteration,acceptance_probability=X_MH_banana$acceptance_rates)
df2<-data.frame(iterations=iteration,acceptance_probability=X_AM_banana$acceptance_rates)
df3<-data.frame(iterations=iteration,acceptance_probability=X_AM2_banana$acceptance_rates)
df4<-data.frame(iterations=iteration,acceptance_probability=X_AM_banana_sh$acceptance_rates)
df5<-data.frame(iterations=iteration,acceptance_probability=X_AM2_banana_sh$acceptance_rates)
df6<-data.frame(iterations=iteration,acceptance_probability=X_AM_banana_th$acceptance_rates)
df7<-data.frame(iterations=iteration,acceptance_probability=X_AM2_banana_th$acceptance_rates)
df8<-data.frame(iterations=iteration,acceptance_probability=rep(0.234,n_iter+1))

ggplot(df1,aes(iterations,acceptance_probability))+geom_line(aes(color="MH"))+
  geom_line(data=df2,aes(color="AM"))+
  geom_line(data=df3,aes(color="AM2"))+
  geom_line(data=df4,aes(color="AM + shrinkage"))+
  geom_line(data=df5,aes(color="AM2 + shrinkage"))+
  geom_line(data=df6,aes(color="AM + thresholding"))+
  geom_line(data=df7,aes(color="AM2 + thresholding"))+
  geom_line(data=df8,aes(color="Optimal acceptance"))+
  labs(color="Alorithm:")

```



*# all lines of the mean in the same plot with a legend*



```

target=pi_norm_corr

X_MH = mcmc(target = target,
            n_iter = n_iter,
            x_1 = x_1,
            adapt=adapt)

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: AM .
## Covariance matrix estimator is: Sample covariance .
## MCMC finished in 0.08897471 seconds.
## The last acceptance rate was: 0.365

X_AM = mcmc(target = target,
            n_iter = n_iter,
            x_1 = x_1,
            adapt=adapt,
            t_adapt = t_adapt,
            cov_estimator=cov_estimator
            )

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: AM .
## Covariance matrix estimator is: Sample covariance .
## MCMC finished in 1.980117 seconds.
## The last acceptance rate was: 0.255

X_AM2 = mcmc(target = target,
            n_iter = n_iter,
            x_1 = x_1,
            adapt="AM2"
            )

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: AM2 .
## Covariance matrix estimator is: Sample covariance .
## MCMC finished in 0.06799889 seconds.
## The last acceptance rate was: 0.35

X_AM_sh= mcmc(target = target,
            n_iter = n_iter,
            x_1 = x_1,
            adapt="AM",
            t_adapt = t_adapt,
            cov_estimator=cov_estimator1
            )

```

```

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: AM .
## Covariance matrix estimator is: Shrinkage estimator .Estimating optimal shrinkage intens
##
##
## MCMC finished in 2.147057 seconds.
## The last acceptance rate was: 0.265

X_AM2_sh = mcmc(target = target,
                n_iter = n_iter,
                x_1 = x_1,
                adapt="AM2",
                cov_estimator=cov_estimator1
                )

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: AM2 .
## Covariance matrix estimator is: Shrinkage estimator .
## MCMC finished in 0.06679249 seconds.
## The last acceptance rate was: 0.38

X_AM_th= mcmc(target = target,
              n_iter = n_iter,
              x_1 = x_1,
              adapt="AM",
              t_adapt = t_adapt,
              cov_estimator=cov_estimator2
              )

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: AM .
## Covariance matrix estimator is: Thresholding estimator .
## MCMC finished in 0.2747104 seconds.
## The last acceptance rate was: 0.315

X_AM2_th = mcmc(target = target,
                n_iter = n_iter,
                x_1 = x_1,
                adapt="AM2",
                cov_estimator=cov_estimator2
                )

## Running MCMC targeting target with 200 iterations.
## Adaptation algorithm is: AM2 .
## Covariance matrix estimator is: Thresholding estimator .
## MCMC finished in 0.07250571 seconds.
## The last acceptance rate was: 0.4

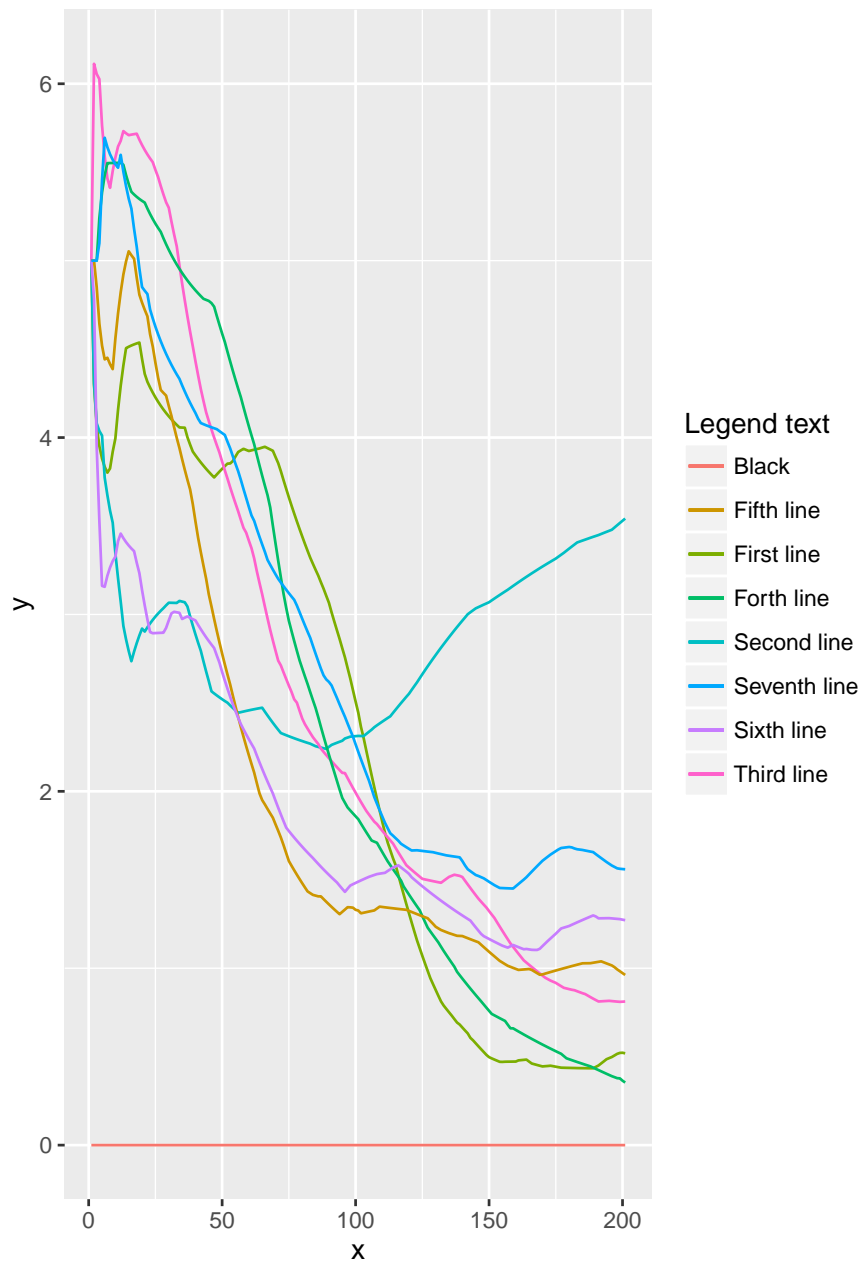
```

```

df1<-data.frame(x=iteration,y=X_MH$sample_mean[,1])
df2<-data.frame(x=iteration,y=X_AM$sample_mean[,1])
df3<-data.frame(x=iteration,y=X_AM2$sample_mean[,1])
df4<-data.frame(x=iteration,y=X_AM_sh$sample_mean[,1])
df5<-data.frame(x=iteration,y=X_AM2_sh$sample_mean[,1])
df6<-data.frame(x=iteration,y=X_AM_th$sample_mean[,1])
df7<-data.frame(x=iteration,y=X_AM2_th$sample_mean[,1])
df8<-data.frame(x=iteration, y=rep(0,n_iter+1))

ggplot(df1,aes(x,y))+geom_line(aes(color="First line"))+
  geom_line(data=df2,aes(color="Second line"))+
  geom_line(data=df3,aes(color="Third line"))+
  geom_line(data=df4,aes(color="Forth line"))+
  geom_line(data=df5,aes(color="Fifth line"))+
  geom_line(data=df6,aes(color="Sixth line"))+
  geom_line(data=df7,aes(color="Seventh line"))+
  geom_line(data=df8,aes(color="Black"))+
  labs(color="Legend text")

```



*# all lines of the mean in the same plot with a legend*

### 3 A Bit About knitr

Whilst this demonstrates that the algorithms both approximately sample from the correct target distribution, there is no improvement in acceptance rate due to the adaptation steps. We require a less regular shaped target distribution to test whether adaptation improves our acceptance rate towards the optimal 0.234. For this we follow Haario et al in using a banana-shaped distribution. The following results show a significantly lower acceptance rate for the adaptive version of the algorithm.

The AM algorithm with banana target:

The MH algorithm with banana target:

We can see that the acceptance probability is closer to the optimal 0.234 with adaptation according to Haario et al.

### 4 Less naive covariance estimation

Naive empirical covariance estimators are unstable for high-dimensional problems with little data; the literature .

Following these two adaptive MH implementations, we are now looking to use something that is less naive than the vanilla estimator for the covariance matrix. In particular we have began testing the shrinkage and thresholding estimators as modifications to AM2. We will now test whether these estimators translate into better convergence by looking at their trajectories. We then plan to include burn in, and compare all algorithms in terms of the suboptimality factor following Roberts and Rosenthal (2009).

### 5 L<sup>A</sup>T<sub>E</sub>X

L<sup>A</sup>T<sub>E</sub>X itself is complicated if you've never used it before, but I'm sure you'll pick it up quickly: there are a lot of guides on the web. I recommend using the `align` environment (in the `amsmath` package) for displayed equations:

$$\begin{aligned} f(x) &= x^3 - x - 1 \\ g(y) &= y^4 + 2y \end{aligned}$$

You can cite in two ways using the `natbib` package: (Haario et al., 2001) and Haario et al. (2001).

### References

A Gelman, GO Roberts, and WR Gilks. Efficient metropolis jumping rules. 1996.

- Heikki Haario, Eero Saksman, and Johanna Tamminen. An adaptive metropolis algorithm. *Bernoulli*, 7(2):223–242, 04 2001. URL <http://projecteuclid.org/euclid.bj/1080222083>.
- Gareth O. Roberts and Jeffrey S. Rosenthal. Examples of adaptive mcmc. *Journal of Computational and Graphical Statistics*, 18(2):349367, 2009. doi: 10.1198/jcgs.2009.06134. URL <http://dx.doi.org/10.1198/jcgs.2009.06134>.