evolMC: a package for Monte-Carlo simulation

W Burchett E Roualdes G Weyenberg

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Getting the Package

Package evolMC.

http://github.com/grady/evol-mc

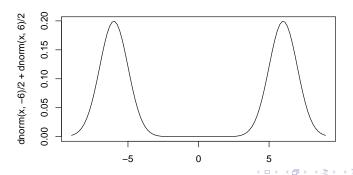
```
library(devtools)
install_github("evol-mc", "grady")
library(evolMC)
```

See the "example" package vignette for demo code.



The Problem

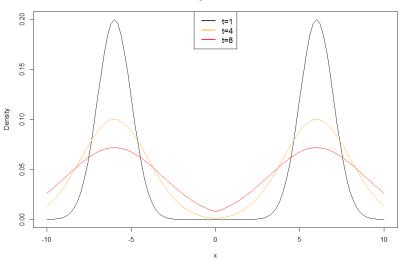
- ▶ We wish to sample from a distribution with multiple modes, separated by regions of very low probability density.
- ► Traditional Metropolis algorithms do not work well here.



Heating a Distribution

- ▶ Consider a distribution $f(x|t) \propto \exp\{-H(x)/t\}$, where $t \geq 1$ is called the *temperature* and H(x), called the *fitness function*, corresponds to the negative log-density of x, up to a constant.
- Increasing the temperature causes the distribution to become "flatter", i.e. to have thicker tails.

Heating a Distribution





Genetic Algorithm Language

- ▶ A population **x** consists of *N* individuals x_i , i = 1, ..., N, where $x_i \in \mathbb{R}^d$.
- Associated with the population is a set of temperatures $\mathbf{t} = (t_1, \dots, t_N)$. It is assumed that the t_i are in descending order, terminating at the temperature of the target distribution. (In most cases this means $t_N = 1$.)
- ▶ Each individual x_i is independently sampled from the distribution $f_i(x_i) \propto f(x_i|t_i)$.
- Parallel Tempering exchanges individuals at different temperatures using a Metropolis-Hastings step to allow the walker to more easily find separated modes.



Algorithm 1 Evolutionary Monte Carlo

procedure EMC

with prob p_m

 \mathbf{Mutate}

otherwise

Crossover

end w/prob

EXCHANGE

end procedure

 $\triangleright p_m$ is the mutation probability.

Algorithm 2 A Gaussian random-walk mutation.

```
procedure MUTATE

Copy the current population to x.

for all individuals x_i in x do

y \leftarrow \mathcal{N}_d(x_i, t_i \sigma^2 I)

with prob \min\{1, \exp(-H(y)/t_i + H(x_i)/t_i)\}

x_i \leftarrow y

end w/prob

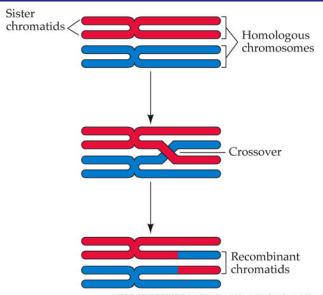
end for

Set current population to x.

end procedure
```

Algorithm 3 The fitness-weighted *crossover*.

```
procedure Crossover
   Copy the current population to x.
   for all individuals x_i in x do
       w_i \leftarrow \exp(-H(x_i))
   end for
   Select k uniformly from \{1: d\}.
   Select i from \{1: N\} with weights proportional to \{w_i\}.
   Select j uniformly from \{1: N\}\setminus\{i\}.
   In x, swap elements k: d of individuals i and j.
   with prob · · ·
                                                           ▶ Metropolis update.
       Set the current population to x.
   end w/prob
end procedure
```







Algorithm 4 The *exchange* attempts to swap individuals between neighboring temperature states.

```
procedure EXCHANGE

Copy the current population to x.

Select i uniformly from \{1: N\}.

Select j uniformly from \{i \pm 1\} \cap \{1: N\}.

Swap individuals i, j of x.

with prob \min\{1, \exp(-H(x_i)/t_i - H(x_j)/t_j + H(x_i)/t_j + H(x_j)/t_i)\}

Set the current population to x.

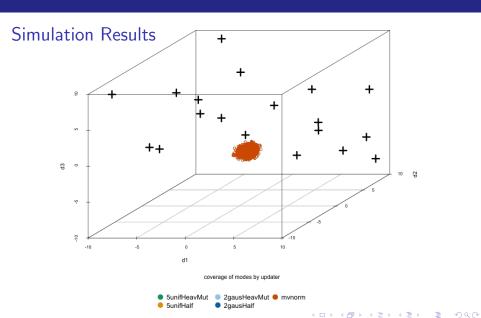
end w/prob

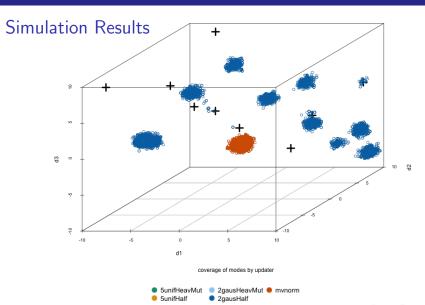
end procedure
```

Simulation Setup

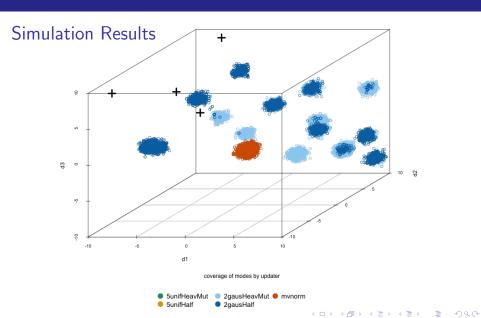
- ▶ Target distribution: 20 part $\mathcal{N}(\mu, 0.1I_3)$ w/ equal mixing proporitions and $\mu \in [-10, 10]^3$.
- ▶ Mutation probabilities: $p_m \in \{0.1, 0.5, 0.9, 1.0\}$
- ▶ Mutation updater: $u_m \in \{uniform, Gaussian\}$
- ▶ Temperatures: $\mathbf{t} \in \{(1,5), (1,2,4,8,15), (1,5,15,30,50)\}$
- MH: multivariate normal proposal distribution
- ▶ Iterations: 2e4

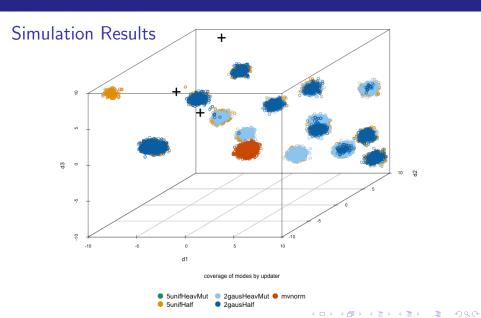


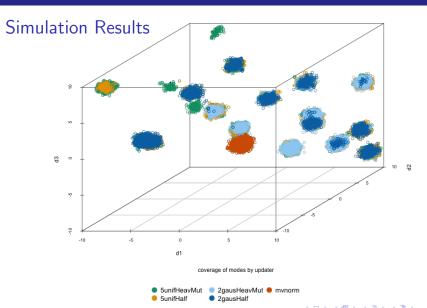












references I

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- W Keith Hastings, Monte carlo sampling methods using markov chains and their applications, Biometrika **57** (1970), no. 1, 97–109.
- Faming Liang, Chuanhai Liu, and Raymond Carroll, Advanced markov chain monte carlo methods: learning from past samples, vol. 714, Wiley. com, 2011.
- Faming Liang and Wing Hung Wong, Evolutionary monte carlo: Applications to cp model sampling and change point problem, Statistica sinica 10 (2000), no. 2, 317–342.

references II



Nicholas Metropolis, Arianna W Rosenbluth, Marshall N Rosenbluth, Augusta H Teller, and Edward Teller, Equation of state calculations by fast computing machines, The journal of chemical physics **21** (1953), 1087.