

Oblique Reflected Diffusion Simulation with Penalty Method

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Introduction

Given a stochastic differential equation (SDE) with oblique reflection :

$$dX(t) = g(X(t))dt + \sigma(X(t))dW(t) + r(X(t))d\ell(t) \quad (1)$$

with $\ell(t)$ is a continuous nondecreasing process with $\ell(0) = 0$ which can increase only when it inside its domain and reflected back inside its domain in the direction of $r(z)$ as X hits the domain at a point z . Once the process is in its domain, the process $X(t)$ behaves as a solution to the SDE below.

$$dX(t) = g(X(t))dt + \sigma(X(t))dW(t). \quad (2)$$

We implement the numerical approximation of solutions of stochastic differential equations driven by brownian motion which are moving inside a domain and obliquely reflected at the boundary as follows:

$$z_{k+1} - z_k = [g(z_k) + \varphi(d(z_k)) \times r(y(z_k))]\epsilon + \sigma(z_k)\epsilon_{k+1}; \quad (3)$$

where $\epsilon_1, \epsilon_2, \dots \sim N(0; \epsilon I_d)$. We define this function Z_ϵ as piecewise constant: $Z_\epsilon(t) := Z_\epsilon(k\epsilon) = z_k, t \in [k\epsilon; (k+1)\epsilon), k = 0, 1, \dots$

Half-line

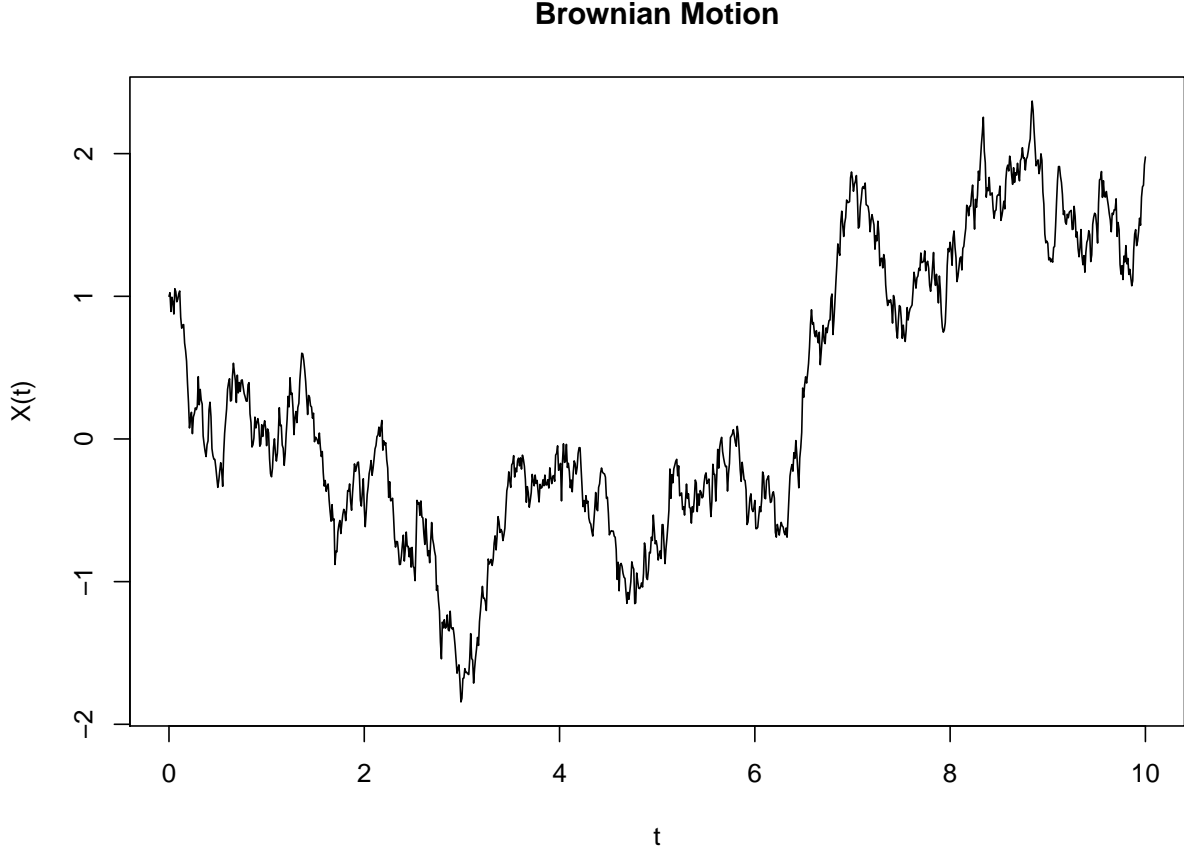
Obliquely reflected brownian motion with penalty term

We simulate a brownian motion with a penalty term for half-line.

$$z(t + \epsilon) \approx z(t) + [\varphi(d(z(t))) \times r(y(z(t)))]\epsilon + \varepsilon_{t+\epsilon},$$

where $Z(0) = 1$, $\varepsilon_{t+\epsilon} \sim \mathcal{N}(0, \epsilon)$, $\epsilon = 0.01$, time horizon $t = 10$ and

$$\varphi(d(z(t))) \times r(y(z(t))) = \text{penalty term with reflection} = \begin{cases} 0, & \text{if } z(t) \geq 0 \\ a[z(t)]^p, & \text{if } z(t) < 0 \end{cases}$$



Convergence on half-line

A reflected brownian motion for unit negative drift has a stationary distribution, which is exponential with mean $\frac{1}{2}$, which serves as a limiting distribution for a long-time limit.

2-D Brownian motion with $\epsilon = 0.01$, $X(0) = 1$ and time horizon $t = 10$

$$Z(\vec{t} + \epsilon) - Z(\vec{t}) \sim \mathcal{N}(\vec{0}, \Sigma)$$

where $\Sigma = \begin{pmatrix} \epsilon & 0 \\ 0 & \epsilon \end{pmatrix}$ and $Z(\vec{.})$ is a BM with 2-D

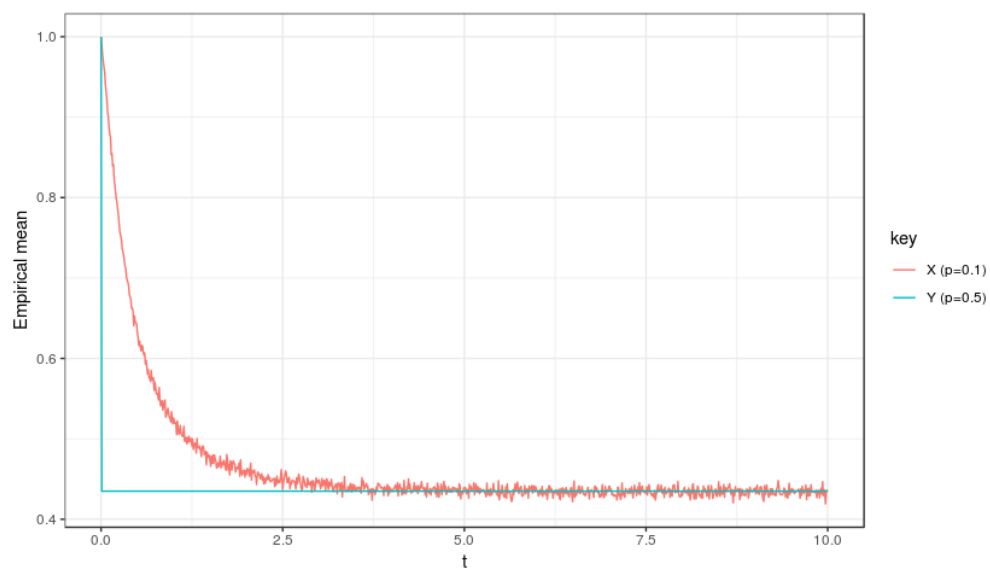
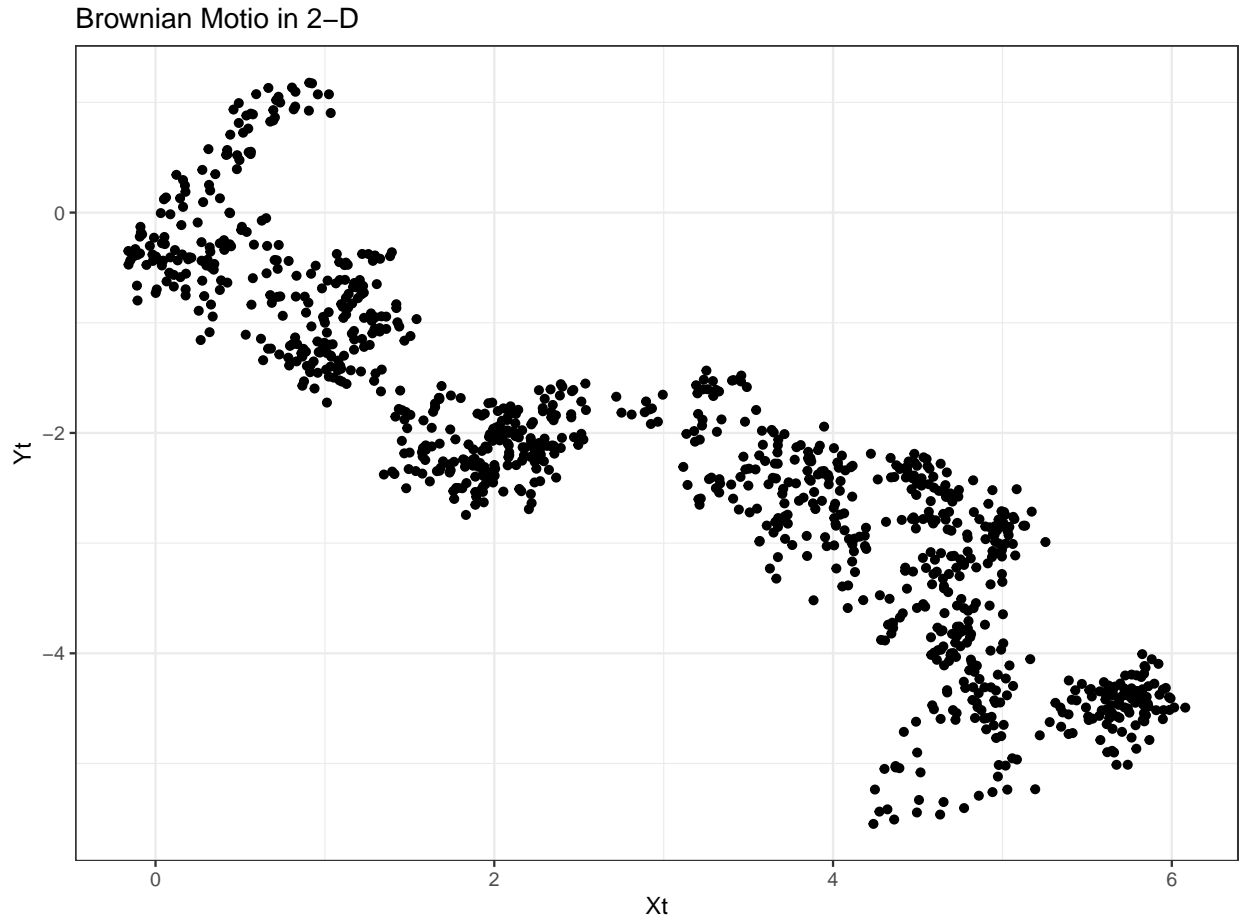


Figure 1: Stationary distribution of reflected brownian motion with unit negative drift

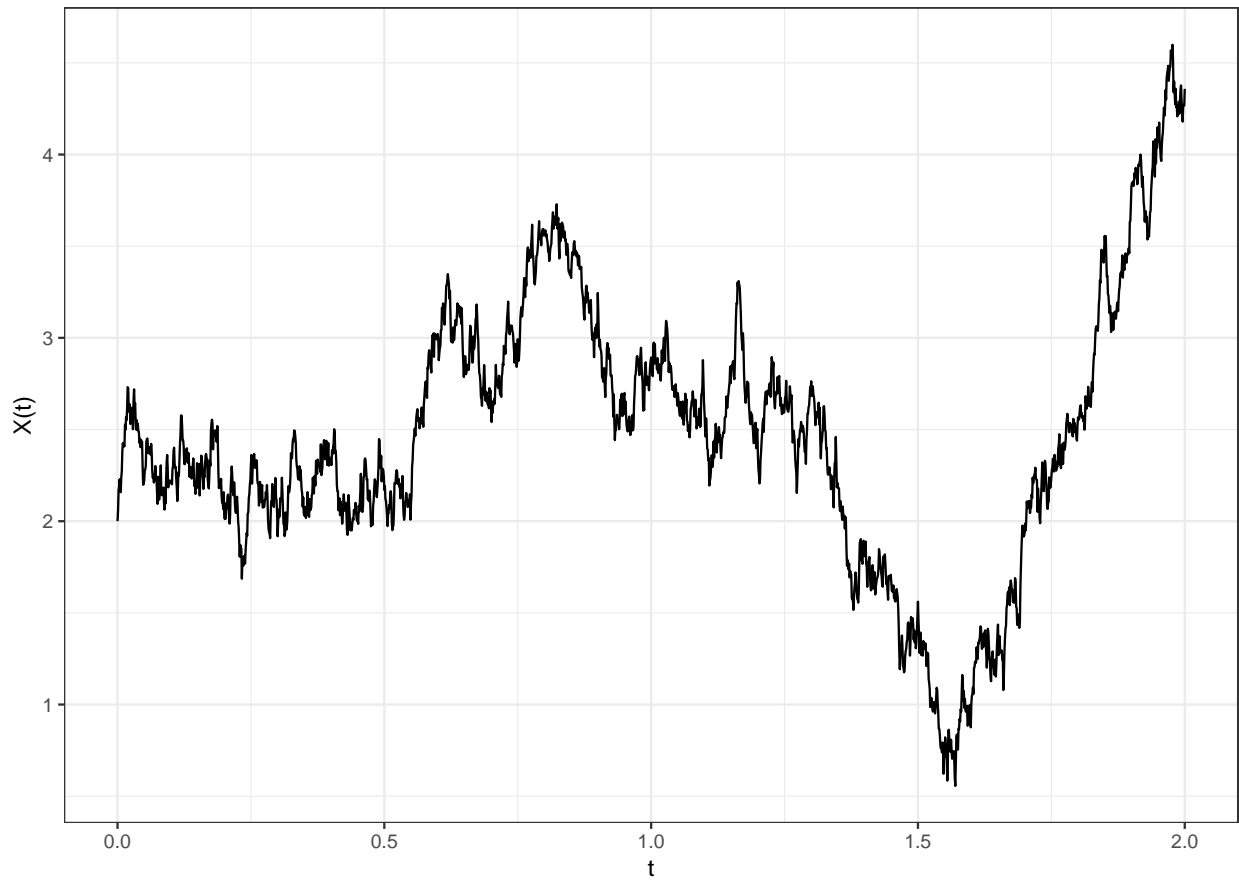




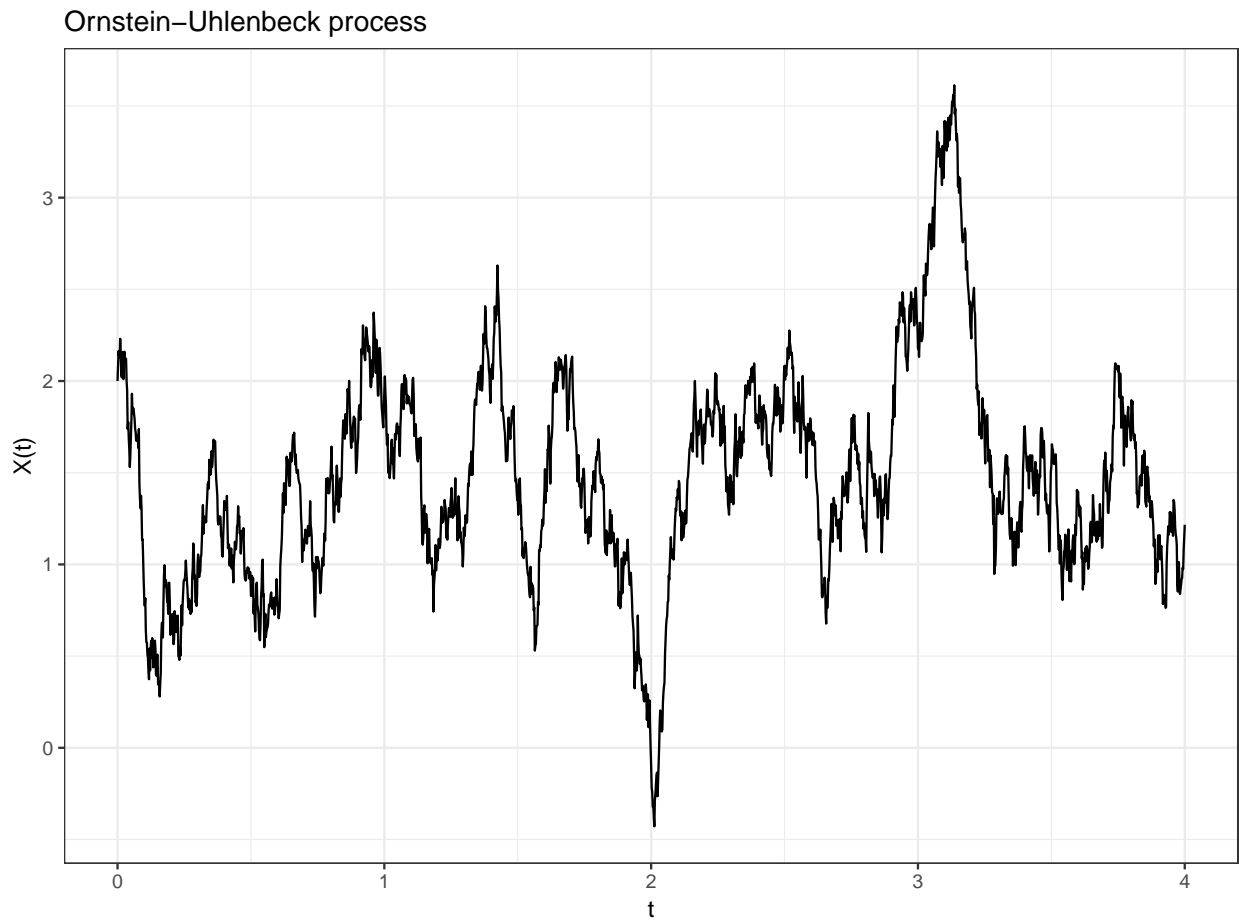
Ornstein-Uhlenbeck process with $\epsilon = 0.001$, $X(0) = 2$ and time horizon $t = 2$

$$dX(t) = 3(2 - X(t))dt + 2dW(t)$$

Ornstein–Uhlenbeck process

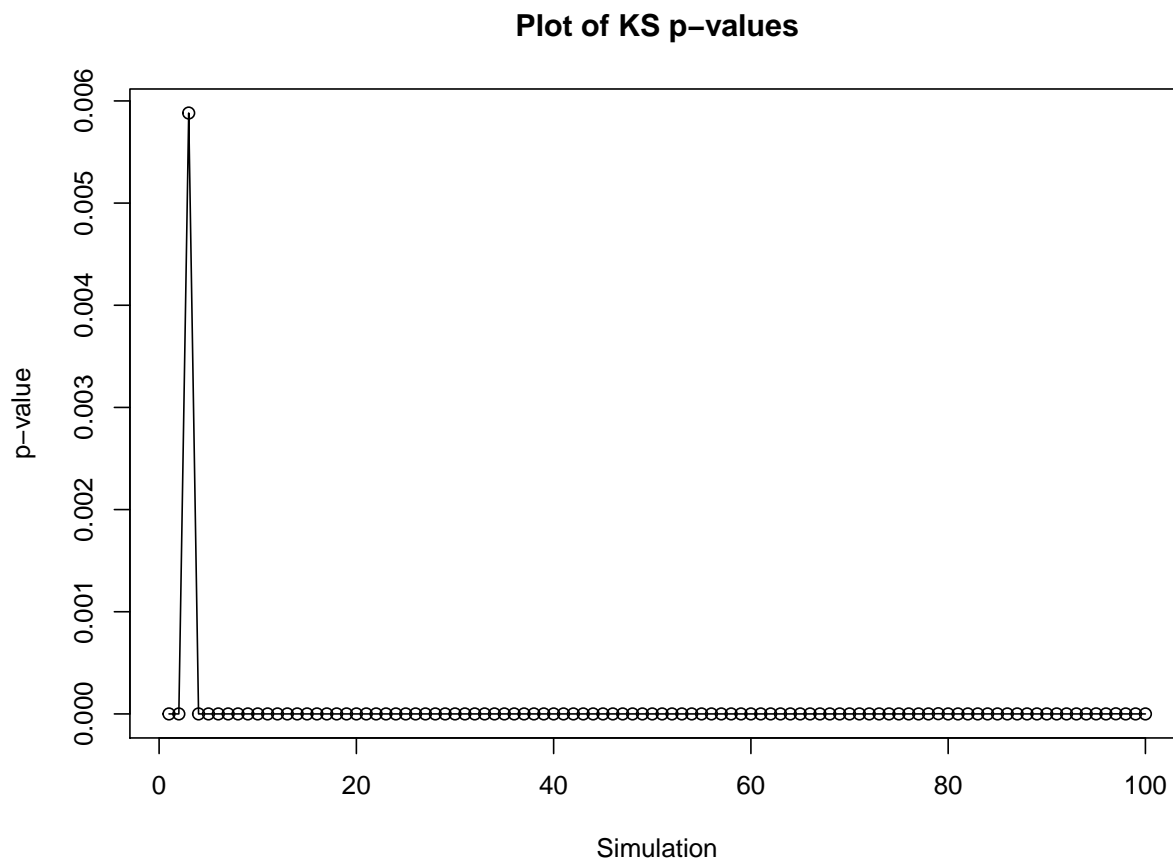


*Ornstein-Uhlenbeck process repeated with step
 $\epsilon = 0.002$, $X(0) = 2$ and time horizon $t = 2$*



[1] 3.589199

Comparing Ornstein-Uhlenbeck processes using KS test



Two-dimensional SDE

Comparing 2-D SDE processes using Peacock test

Convergence on half-line

A reflected brownian motion for unit negative drift has a stationary distribution, which is exponential with mean $\frac{1}{2}$, which serves as a limiting distribution for a long-time limit.

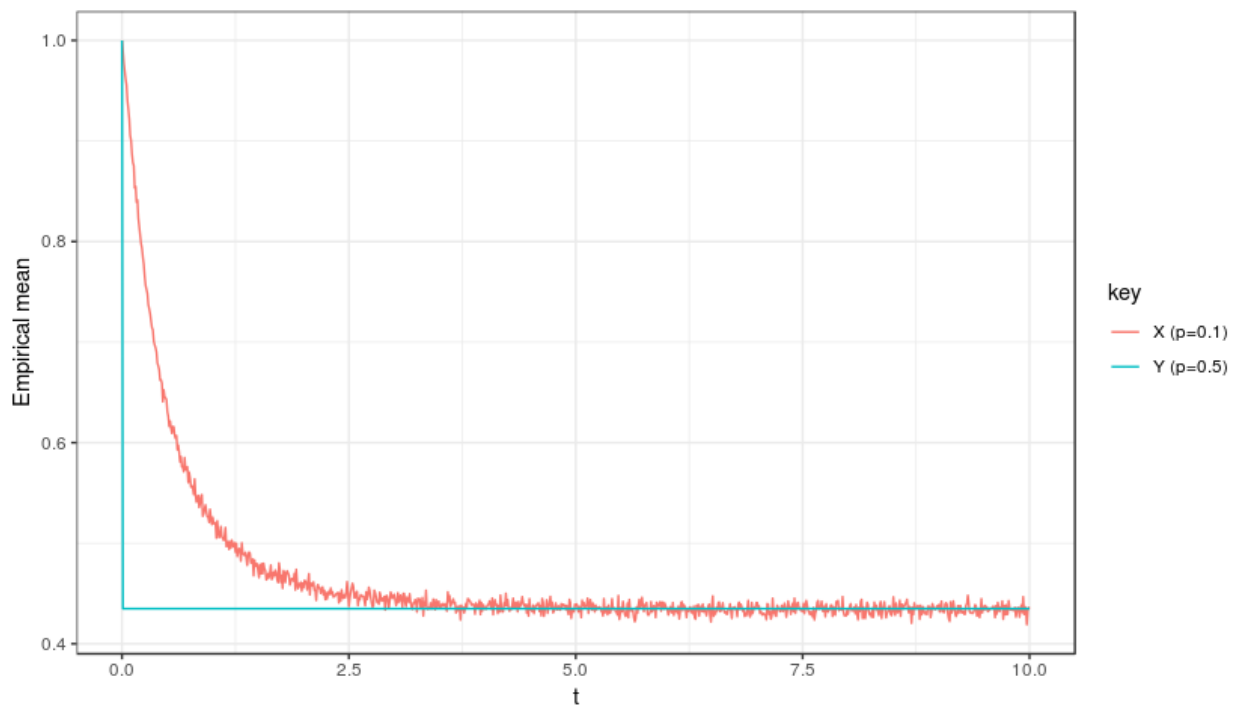


Figure 2: Stationary distribution of reflected brownian motion with unit negative drift

R Code

```
rm(list = ls())
knitr::opts_chunk$set(
  echo = FALSE,      # don't show code
  comment = NA,
  error=FALSE ,
  warning = FALSE,   # don't show warnings
  message = FALSE,   # don't show messages (less serious warnings)
  cache = FALSE,     # set to TRUE to save results from last compilation
  fig.align = "center"# center figures
)

## libraries
library(MASS)
library(dplyr)
library(tidyr)
library(ggplot2)
library(Peacock.test)

### function to simulate standard brownian motion
```



```

BrownianM<-function(Begin, End,Step, X0){
  t<-seq(Begin,End,Step)
  n=length(t)-1
  Xt<-cumsum(c(X0,rnorm(n,0,sqrt(Step))))
  return(data.frame(t,Xt))
}

X1<-BrownianM(0,10,Step = 0.01,X0=1)
plot(X1$t, X1$Xt, ylab = "X(t)",xlab = "t",type="l", lwd=1,
     main="Brownian Motion")

knitr::include_graphics("C:/Users/Charles/Documents/UNR/BOOKS/SEM8/Stochastic Simulation
### function to simulate multidimensional Brownian Motion
Multi_BrownianM<-function(Begin, End,Step, X0){
  t<-seq(Begin,End,Step)
  n=length(t)-1
  mu <- c(0,0) # Mean
  sigma <- matrix(c(Step, 0, 0, Step),2)
  Xt<-apply(rbind(X0, mvrnorm(n, mu = mu, Sigma = sigma )),2,cumsum)
  colnames(Xt) <- c("Xt","Yt")
  return(data.frame(t,Xt))
}

X2<-Multi_BrownianM(0,10,Step = 0.01,X0=c(1,-1))

par(mfrow=c(1,2))
X2%>% gather(key,value, Xt, Yt) %>%
  ggplot(aes(x=t, y=value, colour=key)) +
  geom_line() +theme_bw()+
  ggtitle("Brownian Motion") +
  xlab("t") + ylab("Value")

ggplot(data = X2, aes(x=Xt, y=Yt)) +
  geom_point() +theme_bw()+
  ggtitle("Brownian Motio in 2-D") +
  xlab("Xt") + ylab("Yt")

#plot(X1$Xt,X1$Yt, lwd=2,xlab="X(t)", ylab="Y(t)",
#      main="Brownian Motio in 2-D")

### simulate ORNSTEIN-UHLENBECK Process
#Mu=long run mean, lambda = mean reversion speed sigma=
#set.seed(12345)
ornstein_uhlenbeck <- function(Begin, End,Step,Mu,lambda,sigma,X0){
  t<-seq(Begin,End,Step) ### create sequence

```

```

n=length(t)
dt <- Step
dw<-rnorm(n, 0, sqrt(Step))
X<-NULL
X[1] <- X0
for (i in 2:n) {
  X[i] <- X[i-1] + lambda*(Mu-X[i-1])*dt + sigma*dw[i-1]
}
return(data.frame(t,X));
}

X3<-ornstein_uhlenbeck(0,2,Step = 0.001,Mu=2,lambda = 3,sigma = 2,X0=2)
ggplot(data = X3, aes(x=t, y=X)) +
  geom_line() +theme_bw()+
  ggtitle("Ornstein-Uhlenbeck process") +
  xlab("t") + ylab("X(t)")

X4<-ornstein_uhlenbeck(0,4,Step = 0.002,Mu=2,lambda = 3,sigma = 2,X0=2)
ggplot(data = X4, aes(x=t, y=X)) +
  geom_line() +theme_bw()+
  ggtitle("Ornstein-Uhlenbeck process") +
  xlab("t") + ylab("X(t)")

### Quality of simulation
Distance<- max(abs(X4-X3))
Distance
ks_p_value<-NULL
K<-100
for (i in 1:K) {
  Yt1<-ornstein_uhlenbeck(0,2,Step = 0.001,Mu=2,lambda = 3,sigma = 2,X0=5)
  Yt2<-ornstein_uhlenbeck(0,4,Step = 0.002,Mu=2,lambda = 3,sigma = 2,X0=5)
  ks_p_value[i]<-ks.test(Yt1$X,Yt2$X)$p.value
}

plot(ks_p_value,type="o", xlab="Simulation",ylab="p-value",
     main="Plot of KS p-values")
points(rep(0.05,K),type = "l", col="red")

SDE_2D <- function(Begin, End,Step,X0){
  mu <- c(0,0) # Mean
  sigma <- matrix(c(Step, 0, 0, Step),2)
  t<-seq(Begin,End,Step) ### create sequence
  n=length(t)
  dt <- Step

```

```

X<- matrix(NA, nrow = n, ncol = 2)
X[1,] <- X0
for (i in 2:n) {
  drf1<-(t(c(1,-1,1))%%matrix(c(X[(i-1), ],1)))*dt
  drf2<- ((t(c(-2,-1))%%matrix(X[(i-1), ])))*dt
  Wt<- c(2*X[(i-1),1]-3,4)* mvrnorm(1, mu = mu, Sigma = sigma )
  X[i,] <- c(X[(i-1), ]) + c(drf1,drf2) + Wt
}
X<-data.frame(X)
colnames(X)<-c("X1t", "X2t")
return(data.frame(t,X))
}
X6<-SDE_2D(0,2,Step = 0.001,X0=c(1,-3))
X6%>% gather(key,value, X1t, X2t) %>%
  ggplot(aes(x=t, y=value, colour=key)) +
  geom_line() +theme_bw()+
  ggtitle("SDE") +
  xlab("t") + ylab("Value")
ggplot(data = X6, aes(x=X1t, y=X2t)) +
  geom_point() +theme_bw()+
  ggtitle("SDE in 2-D") +
  xlab("X1(t)") + ylab("X2(t)")

peacock_p_value<-NULL
K<-100
for (i in 1:K) {
  Yt7<-SDE_2D(0,2,Step = 0.001,X0=c(1,-3))
  Yt8<-SDE_2D(0,4,Step = 0.002,X0=c(1,-3))
  peacock_p_value[i]<-peacock2(Yt8[,2:3],Yt7[,2:3])
}

plot(peacock_p_value,type="o", xlab="iteration",ylab="p-value",
      main="Plot of KS p-values")
points(rep(0.05,K),type = "l", col="red")

knitr::include_graphics("C:/Users/Charles/Documents/UNR/BOOKS/SEM8/Stochastic Simulation
# this R markdown chunk generates a code appendix

```