

Computational Statistics

Lab 6

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2017-03-08

Question 1

1.1

```
genfunc <- function(x) {  
  (x^2 / exp(x)) - 2 * exp(-(9 * sin(x)) / (x^2 + x + 1))  
}
```

1.2

```
crossover <- function(x,y) {  
  (x + y) / 2  
}
```

1.3

```
mutate <- function(x) {  
  x^2 %% 30  
}
```

1.4

```
genetic <- function(maxiter, mutprob) {  
  ## a)  
  ## plot(x = 0:30, y= genfunc(0:30), xlim = c(0,30), type = "l", xlab="", ylab="")  
  
  ## b)  
  X <- seq(0,30,by = 5)  
  
  ## c)  
  values <- genfunc(X)  
  ## points(X, values, col = "red")  
  
  ## d)  
  bestvalue <- -Inf  
  
  for (i in 1:maxiter){  
  
    ## i
```

```

    parents <- sample(1:length(X), size = 2 )

    ## ii
    victim <- which.min(values)

    ## iii
    child <- crossover(X[parents[1]],X[parents[2]])

    if (mutprob > runif(1,0,1)) {
      child <- mutate(child)
    }

    ## iv
    X[victim] <- child

    values[victim]<- genfunc(child)
    ## values <- genfunc(X)

    ## v
    bestvalue <- max(bestvalue, max(values))
  }

  ## points(x = X, y = values, col = "darkgreen")
  list(opt=bestvalue, pop=X, vals=values)
}

```

1.5

```

func_data <- data.frame(x=seq(0, 30, by=0.1), y=genfunc(seq(0, 30, by=0.1)))

set.seed(123456)
r1 <- genetic(maxiter = 10, mutprob = 0.1)
r1$opt

## [1] -1.7

set.seed(123456)
r2 <- genetic(maxiter = 10, mutprob = 0.5)
r2$opt

## [1] 0.14

set.seed(123456)
r3 <- genetic(maxiter = 10, mutprob = 0.9)
r3$opt

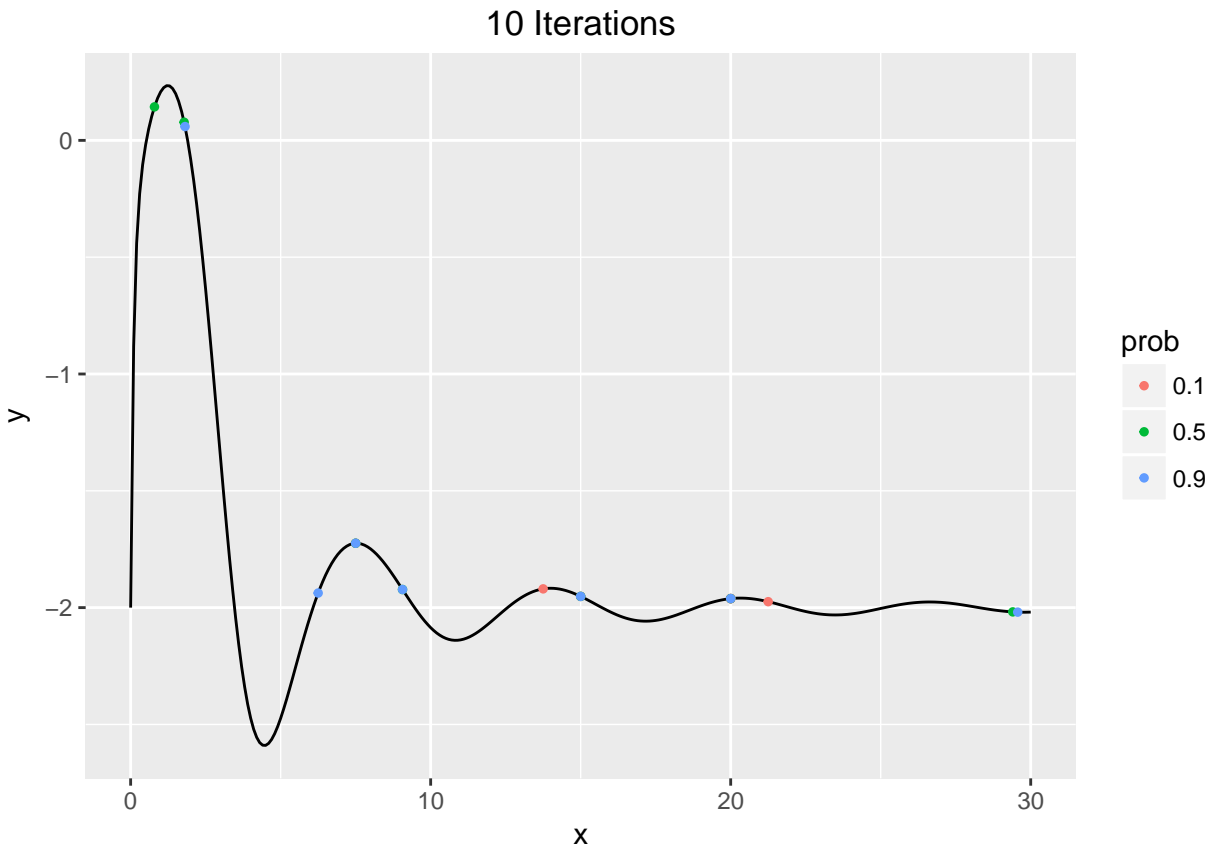
## [1] 0.059

rd1 <- data.frame(x=r1$pop, y=r1$vals, prob="0.1")
rd2 <- data.frame(x=r2$pop, y=r2$vals, prob="0.5")
rd3 <- data.frame(x=r3$pop, y=r3$vals, prob="0.9")

plot_data <- rbind(rd1, rd2, rd3)

```

```
ggplot() +
  ggtitle("10 Iterations") +
  geom_line(data=func_data, aes(x=x, y=y)) +
  geom_point(data=plot_data, aes(x=x, y=y, col=prob), size=1) +
  theme(plot.title=element_text(hjust=0.5))
```



```
set.seed(123456)
r1 <- genetic(maxiter = 100, mutprob = 0.1)
r1$opt
```

```
## [1] -1.7
```

```
set.seed(123456)
r2 <- genetic(maxiter = 100, mutprob = 0.5)
r2$opt
```

```
## [1] 0.23
```

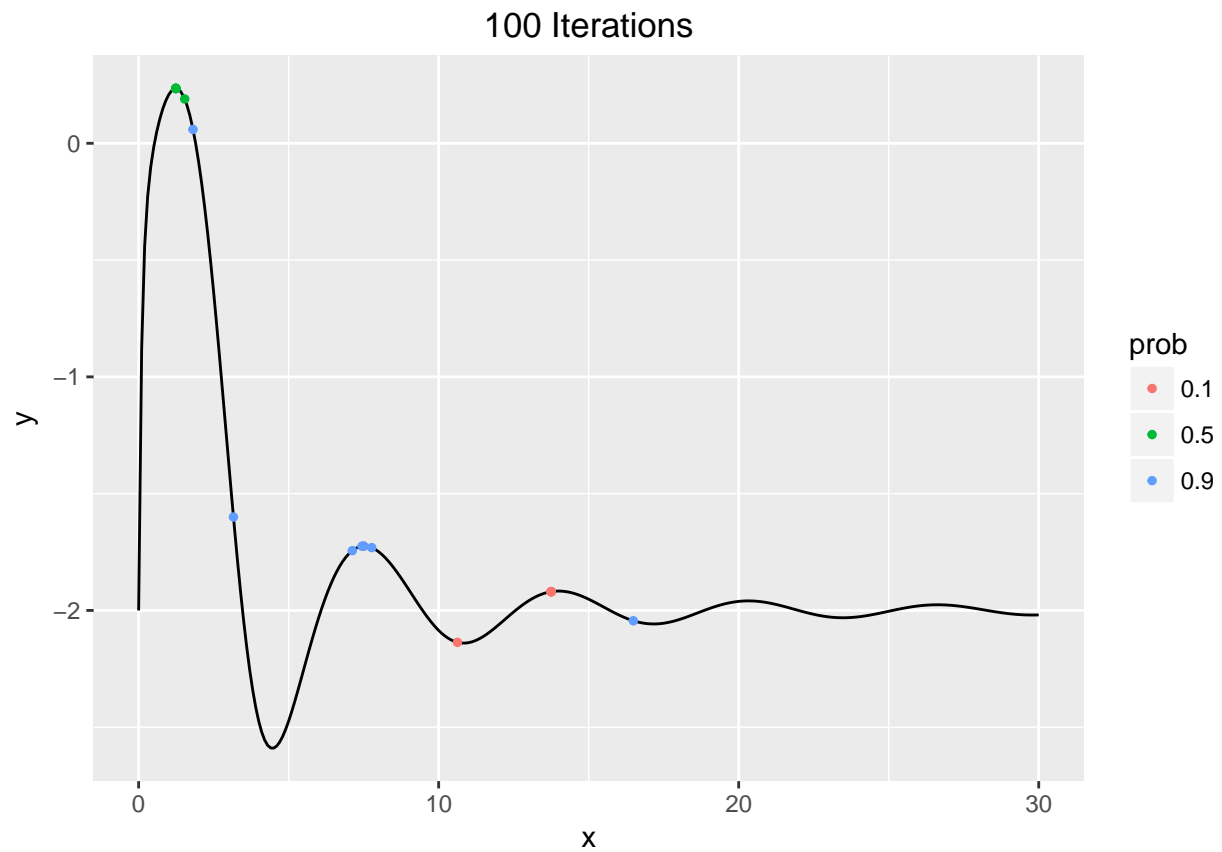
```
set.seed(123456)
r3 <- genetic(maxiter = 100, mutprob = 0.9)
r3$opt
```

```
## [1] 0.059
```

```
rd1 <- data.frame(x=r1$pop, y=r1$vals, prob="0.1")
rd2 <- data.frame(x=r2$pop, y=r2$vals, prob="0.5")
rd3 <- data.frame(x=r3$pop, y=r3$vals, prob="0.9")
```

```
plot_data <- rbind(rd1, rd2, rd3)

ggplot() +
  ggtitle("100 Iterations") +
  geom_line(data=func_data, aes(x=x, y=y)) +
  geom_point(data=plot_data, aes(x=x, y=y, col=prob), size=1) +
  theme(plot.title=element_text(hjust=0.5))
```



Question 2

2.1

2.2

2.3

2.4