HW 5 Random Number Generators

STAT 5400

Due: Oct 4, 2024 9:30 AM

Problems

Submit your solutions as an .Rmd file and accompanying .pdf file. Include all the **relevant** R code and output. Always interpret your result whenever it is necessary.

Problems

1. Generators of exponential distributions

• Fill in the code on Slide 36 of S3P1.pdf. You may either use the pdf of exp(2) manually or call dexp in R.

```
set.seed(5400)
Ulist <- runif(1000, 0, 1)
Xlist <- -2 * log(Ulist)

par(mfrow = c(1,2))
hist(Xlist)
plot(seq(0.001, 12, len=200), dexp(seq(0.001, 12, len=200), rate = 1, log = FALSE), type="l", ylim = c(</pre>
```

Histogram of Xlist exp(2)300 0.8 Frequency 9.0 200 100 0.2 0.0

• Fill in the code on Slide 37 of S3P1.pdf.

5

10

Xlist

15

0

0

• (Negative exponential distribution) Suppose there is a distribution with pdf $\frac{1}{2}exp\{\frac{1}{2}(x)\}$, x < 0. Generate a random sample with 1000 observations from such distribution. Have a Q-Q plot to check whether the sample comforts with this distribution.

0

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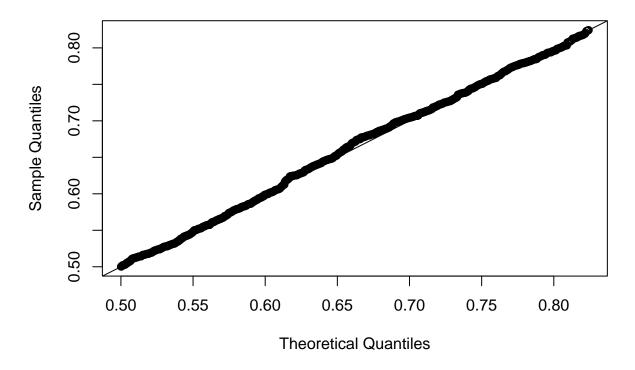
Χ

8

10

```
ulist <- runif(1000, 0, 1)
xlist \leftarrow 0.5 * exp(0.5 * ulist)
theoretical_quantiles <- 0.5 * exp(0.5 * qunif(ppoints(1000)))
plot(theoretical_quantiles, sort(xlist),
xlab="Theoretical Quantiles", ylab="Sample Quantiles",
main="Q-Q plot for exponential distribution")
abline(0, 1)
```

Q-Q plot for exponential distribution



2. Generators of Cauchy distributions.

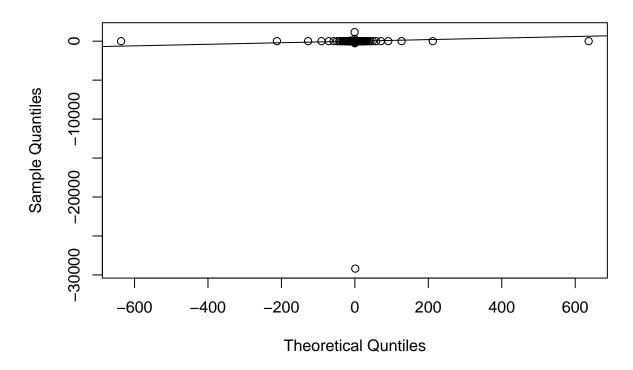
- Implement an algorithm to generate the standard Cauchy distribution using the inverse CDF.
- Check if the generated numbers conform to the standard Cauchy distribution. Basically, you may check this by filling in the code on Slide 42 of S3P1.pdf.

```
u <- runif(1000)

cauchy_samples <- tan(pi * (u - 0.5))

probs = ppoints(1000)
plot(qcauchy(ppoints(1000)), cauchy_samples,
xlab="Theoretical Quntiles",
ylab="Sample Quantiles",
main="Q-Q plot for Cauchy distribution")
abline(0, 1)</pre>
```

Q-Q plot for Cauchy distribution



```
dev.off()
```

```
## null device
## 1
```

3. Generators of Beta distributions and t distributions.

• Generate a Beta(α, α) distribution using

$$X = \frac{1}{2} + \frac{\mathbb{I}_{U_3 \leqslant 1/2}}{2\sqrt{1 + \frac{1}{\left(U_1^{-1/\alpha} - 1\right)\cos^2(2\pi U_2)}}}.$$

where U_1 , U_2 , and U_3 are independently generated random vaiables on (0,1), and the indicator function $\mathbb{I}_{U_3 \leq 1/2}$ takes the value of 1 if the condition is true or the value of -1 otherwise.

```
beta <- function(a,b){
  ind_u3 = ifelse(runif(1)> 0.5,1,-1)

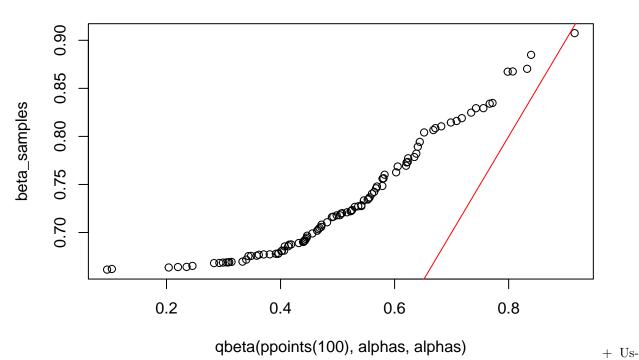
return (0.5 + ind_u3/(2* sqrt(1+1/(((runif(1)^(-1/a))-1)* cos(2 * pi * runif(1))^2) )))
}
```

• With several values of α , generate a sample of 100 observations from Beta(α , α). Have a Q-Q plot to check whether the sample comforts with the beta distribution. You may use the built-in function in R to give the inverse CDF function.

```
set.seed(123)
alphas <- runif(100, 1, 10) # Randomly generate 100 different alpha values between 1 and 10
beta_samples <- beta(alphas, alphas)

qqplot(qbeta(ppoints(100), alphas, alphas), beta_samples, main="Q-Q plot for 100 samples with different abline(0, 1, col="red")</pre>
```

Q-Q plot for 100 samples with different alphas



ing the above method to generate $Y \sim \text{Beta}(n,n)$. Generate a t-distribution with degree of freedom 2n using

$$Z = \frac{\sqrt{n}(Y - 1/2)}{2\sqrt{Y(1 - Y)}}.$$

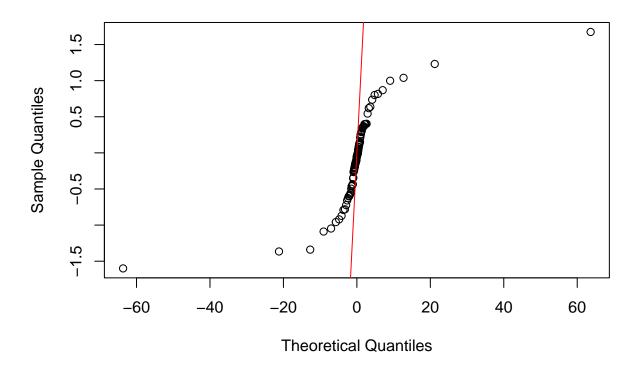
```
generate_t_distribution <- function(n, sample_size = 100) {
  beta_samples <- replicate(sample_size, beta(n)) # Generate Beta(n, n) samples

# Apply the given formula to generate t-distributed samples
  t_samples <- sqrt(n) * (beta_samples - 0.5) / (2 * sqrt(beta_samples * (1 - beta_samples)))
  return(t_samples)
}</pre>
```

• Generate a sample of 100 observations from t(1), namely Cauchy distribution. Have a Q-Q plot to check whether the sample comforts with the t(1) distribution. Compare the Q-Q plot with the one you plotted in the previous question.

```
t_samples <- generate_t_distribution(1)
qqplot(qt(ppoints(100), df=1), t_samples,</pre>
```

Q-Q Plot for t-distribution with df = 1



4. More on accept-reject sampling.

• Generate n values from the truncated normal distribution:

$$Y_i \sim (X|a < X < b)$$
, where $X \sim N(\mu, \sigma^2)$.

Hint: generate observations from normal distribution using 'rnorm" and discard it if it falls outside the interval.

This function should have five arguments:

- n: number of observations,
- mu: the value of mean μ ,
- sigma: the value of standard deviation σ ,
- a: the left endpoint of the interval of -Inf,
- b: the right endpoint of the interval of Inf, This function should return a vector of n truncated normal variables.

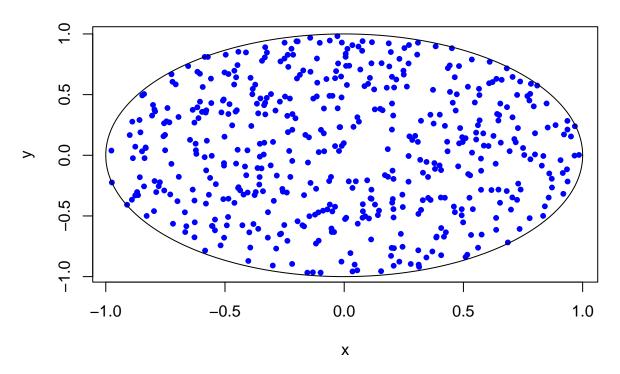
```
} else {
    i = i + 1
    vec.append(val)
    }
}
return(vec)
}
```

5. Uniformly generating points within a circle.

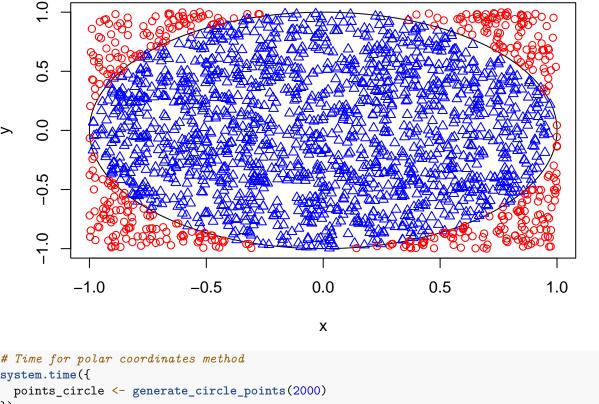
- Implement an algorithm to perform the following steps to uniformly generating points within a circle.
 - Generate a random angle θ uniformly distributed in the range $[0, 2\pi)$.
 - Genreate a random radius r uniformly distributed in the range [0, 1) with a square root scale, i.e., $\mathbf{r} \leftarrow \operatorname{sqrt}(\operatorname{runif}(1))$.
 - Give polar coordinates (r,θ) , get the Cartesian coordinates $x=r\cos\theta$ and $y=r\sin\theta$.
- Generate 500 points and use the code on Slide 48 of S3P1.pdf to plot the points.
- Compare the run time between this algorithm and the one based on accept/reject sampling (Slide 45 of S3P1.pdf).

```
# Function to generate uniformly distributed points within a circle
generate_circle_points <- function(n) {</pre>
  theta <- runif(n, 0, 2 * pi) # Generate random angles between 0 and 2pi
  r <- sqrt(runif(n))
                                   # Generate random radii with square root scaling
                                   # Convert to Cartesian coordinates (x, y)
  x \leftarrow r * cos(theta)
 y <- r * sin(theta)
 return(cbind(x, y))
                                   # Return as matrix of x and y values
}
# Set seed for reproducibility
set.seed(5400)
# Generate 500 points
points_circle <- generate_circle_points(500)</pre>
# Plot the points within the circle
plot(points_circle, pch=20, col="blue", xlab="x", ylab="y", main="Uniform Points in a Circle")
theta <- seq(0, 2 * pi, length.out = 1000)
lines(cos(theta), sin(theta)) # Draw the unit circle boundary
```

Uniform Points in a Circle



Accept/Reject Sampling



```
# Time for polar coordinates method
system.time({
})
##
            system elapsed
      user
          0
##
                   0
# Time for accept/reject method
system.time({
  dat <- matrix(runif(n * 2, -1, 1), n, 2)</pre>
  accept \leftarrow (dat[, 1]^2 + dat[, 2]^2) \leftarrow 1
  accepted_points <- dat[accept, ]</pre>
})
##
      user
            system elapsed
```

6. (Optional) Advanced readings on accept-reject sampling. Read more on accept-reject sampling on Section 4.7 and 4.8 of http://statweb.stanford.edu/~owen/mc/Ch-nonunifrng.pdf. Implement Algorithm 4.8, which generates a gamma distribution using the accept-reject sampling.

##

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