

# Q3 - Rejection Sampling

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## 1 Q3 - Rejection Sampling

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### Important Notes

- The *target* distribution's density function will be denoted with  $f(x)$
- The *trial* distribution's density function will be denoted with  $g(x)$

### Basic Algorithm

1. Generate  $U \sim \text{Uniform}(0, 1)$
2. Generate  $X \sim g(x)$
3. If  $U \leq \frac{f(X)}{M \cdot g(X)}$  then accept  $X$  as a realization from  $f(x)$ , otherwise reject  $X$  and try again

### Goals

Our goal in this notebook is to generate random samples for the Normal distribution using the Cauchy distribution as our trial.

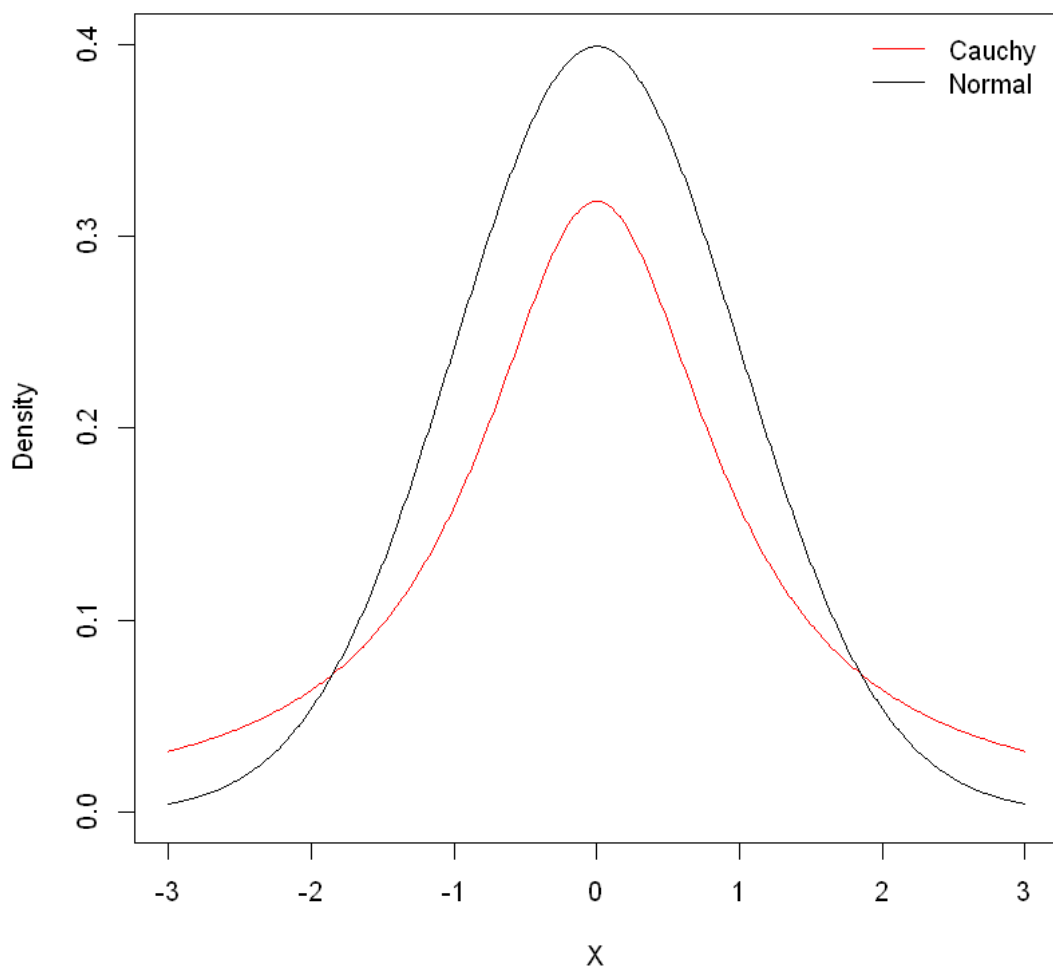
### 1.1 Step 1: Visualizing Our Distributions

Here is a quick visualization so things can stand in perspective.

```
[1]: x <- seq(-3, 3, .01)

plot(x, dcauchy(x), type="l", col="red", ylim=c(0, .4), xlab="X",
      ylab="Density")
lines(x, dnorm(x))

legend("topright", legend=c("Cauchy", "Normal"), lty=1, col=c("red", "black"),
      bty="n")
```



## 1.2 Step 2: Calculating $M$

$$f(x) = \frac{1}{\sqrt{2\pi}} e^{\frac{-x^2}{2}}$$

$$g(x) = \frac{1}{\pi} \frac{1}{1+x^2}$$

$$\frac{f(x)}{g(x)} = \sqrt{\frac{\pi}{2}} (1+x^2) e^{\frac{-x^2}{2}} \leq \sqrt{\frac{2\pi}{e}} \simeq 1.52$$

Please take note that we used the following piece of code from Python's **SymPy** to calculate the limits of  $\frac{f(x)}{g(x)}$ :

```

from sympy import *
init_session()

# define functions
f = 1/sqrt(2*pi) * exp(-x**2 / 2)
g = 1/pi * 1/(1 + x**2)

expr = f/g

# calculate derivative
diff_expr = diff(expr, x)

# solve for "diff_expr = 0"
answers = solveset(diff_expr)
print(answers)  # {-1, 0, 1}

# plot function so we can determine minima and maxima
plot(expr)  # 0 returns a minima and -1, 1 are global maximas

# the limit of "expr"
answer = expr.subs(x, -1).evalf()
print(f"The limit is {answer:.2f}")

```

### 1.3 Step 3: Coding it All

```

[2]: # sample size
n <- 10000
M <- 1.52

# function to determine acception/rejection
accept <- function(x) {
  u <- runif(1)
  if ( (1/M * (pi / 2)^.5 * (1 + x^2) * exp(-x^2 / 2)) > u) {
    return(TRUE)
  }
  return(FALSE)
}

# trials
X <- rep(0, n)
niter <- 0
sample <- NULL

for (i in 1:n) {
  # keep trying until acception
  flag <- FALSE

```

```

while (flag == FALSE) {
  # increment counter
  niter <- niter + 1

  # generate candidate
  sample <- rcauchy(1)
  flag <- accept(sample)
}

X[i] <- sample
}

```

```
[3]: mean(X)
```

```
0.00108225520591382
```

```
[4]: var(X)
```

```
0.980452787713117
```

```
[5]: niter
```

```
15280
```

We can see that the mean and variance are very close to the theoretical mean and variance of the Normal distribution. Also note that the number of iterations required is

$$M \cdot n = 1.52 \times 10000 \simeq 15200$$

Also, keep in mind that

$$P(\text{accept}) = \frac{1}{M} = \frac{1}{52} = .66$$

## 1.4 Step 4: Generating 5 Random Variates

```

[6]: n <- 5

X <- rep(0, n)
sample <- NULL

for (i in 1:n) {
  # keep trying until acception
  flag <- FALSE
  while (flag == FALSE) {
    # generate candidate
    sample <- rcauchy(1)
    flag <- accept(sample)
  }
}

```

```
    X[i] <- sample  
  }
```

```
print(X)
```

```
[1] 0.05767254 0.03340663 0.03644511 1.49799635 -1.25518305
```