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# The Accept-Reject Method: Anthony Gusman, Nicholas Sullivan

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## Introduction

The process of generating pseudo random numbers using a given density function of a random variable is easy if the inverse of the CDF is known or quick to find. However, the general method breaks down when there does not exist a happy way to find the inverse of the CDF. This is why we will be exploring something called the *Acceptance-Rejection Method*.

## Overview of Method

The Acceptance-Rejection Method is actually quite simple. We implement the method with our `acceptReject` function as described in plain English below:

1. Given a probability density function (**PDF**)  $f_X$  of a random variable  $X$  with compact support find its maximum over its range (call it  $c$ ).
2. Generate a pair of random numbers  $(X, Y)$  such that  $X \sim U[\text{support of } f_X]$  and  $Y \sim U[0, c]$ .
3. For each point  $(x_i, y_i)$  if  $0 \leq y_i \leq f_X(x_i)$  then it is "accepted" otherwise it is "rejected."

As you can see this is essentially the same process used in Monte-Carlo integration.

## The Functions

We provide two functions in this paper, `acceptReject.m` and `acceptRejectPlot.m` with the following outputs and arguments:

`[X,Y] = acceptReject(f,a,b,n)`

- **Output:** `[X,Y]`, random numbers defined as above split into a cell with the first row as the accepted values and the second row as the rejected values
- **Input:** `f`, PDF function  $f_X$

- a, lower bound of f's range
- b, upper bound of f's range
- n, number of  $(x_i, y_i)$  pairs (increase this for higher accuracy)

acceptRejectPlot(fun,a,b,Xc,Yc)

- **Output:** plots, one of the pdf with a histogram of the accepted number generated and one of the pdf along with all accepted and rejected points.
- **Input:** fun, PDF function  $f(x)$
- a, lower bound of fun's range
- b, upper bound of fun's range
- Xc, the X returned by acceptReject
- Yc, the Y returned by acceptReject

## Accept Reject and Plotter function code

here is the full code of both functions with comments and syntax high lighting: **acceptReject.m**

[include>acceptReject.m</include](#)

**acceptRejectPlot.m**

[include>acceptRejectPlot.m</include](#)

## Examples

Lets do the beta and someother unhappy distribution

## Full function code

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