

# Stat 243 Final Project Writeup

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## Functions

We created a very modular solution for our Adaptive-Rejection Sampler, which handled unique subtasks within individual functions. Below, we list each function we created and its modular purpose:

### **get\_initial\_abscissae()**

This function builds the vector of initial abscissae. If both endpoints of the inputted domain are finite, then the function simply uses those endpoints (plus/minus a small term to protect against closed intervals) as two of the abscissae and samples the rest of the abscissae uniformly on the domain. If either of the endpoints is infinite, then the function iterates over values until it finds one whose first derivative is of the appropriate sign. For example, if  $D$  is unbounded on the right, we began at  $x = 2$  and keep moving right until we find an  $x$  for which the derivative is positive. Again, once we have two finite abscissae based on the domain endpoints, we sample uniformly on the interval bounded by them to select the rest abscissae.

This function is called at the beginning of `ars()` to set the initial abscissae.

### **get\_z() and get\_z\_all()**

Together, these functions use the abscissae and the log density to find the vector of points at which the tangent lines at the abscissae intersect. They utilize the equations presented in the Gilks, et. al. paper to calculate the vector of  $z$ 's. The `get_z_all()` function is called within `ars()` to recreate the  $z$  vector each time the abscissae get updated.

### **get\_u\_segment() and get\_u()**

Together, these functions use the abscissae and the log density to build the piecewise upper bound of the log density using tangent lines at  $x$ . Again, these functions utilize the formulas from Gilks, et. al.

We represent this piecewise function as a list of slopes and intercepts rather than as a closure-type variable in order to more easily calculate integrals analytically in later functions. The `get_u()` function is called within `ars()` to recreate the upper bound each time the abscissae get updated.

### **get\_l\_segment() and get\_l()**

These functions work very similarly to `get_u_segment()` and `get_u()`, but instead build the piecewise lower bound by calculating the slopes and intercepts of the chords between adjacent abscissae, again using the formulas presented in Gilks, et. al.

Again, we represent this piecewise function as a list of slopes and intercepts. The `get_l()` function is called within `ars()` to recreate the lower bound each time the abscissae get updated.

### **get\_s\_integral()**

This function calculates the integrals under each piecewise element of  $s$ , which is the normalized exponential of the  $u$  function. The function works by calculating the integral under each piecewise element of  $s$  analytically, using the formula as calculated below

$$\int_{z_1}^{z_2} \exp(u_j(t))dt = \int_{z_1}^{z_2} \exp(a_j + b_j t)dt = \int_{a_j + b_j z_1}^{a_j + b_j z_2} \exp(y) \frac{1}{b_j} dy = \frac{1}{b_j} [\exp(y)]_{a_j + b_j z_1}^{a_j + b_j z_2} = \frac{\exp(a_j + b_j z_2) - \exp(a_j + b_j z_1)}{b_j}$$

where  $a_j$  and  $b_j$  are the intercept and slope of the  $j^{th}$  segment of  $u$  respectively. The function returns a vector of integrals, of which the  $j^{th}$  element corresponds to the integral under the  $j^{th}$  segment of  $s$ .

### **get\_l\_integral()**

This function calculates the integrals under each piecewise element of  $l$ , which are piecewise lower hulls underneath the original density function. The function returns a vector of integrals from  $x \in \{x_j, x_{j+1}\}$  for  $j = 1, \dots, k - 1$ , where  $k$  is the subset of  $D$  under consideration. This is performed analytically as follows:

$$\int_{x_1}^{x_2} \exp(l_j(t))dt = \int_{x_1}^{x_2} \exp(a_j + b_j t)dt = \int_{a_j + b_j x_1}^{a_j + b_j x_2} \exp(y) \frac{1}{b_j} dy = \frac{1}{b_j} [\exp(y)]_{a_j + b_j x_1}^{a_j + b_j x_2} = \frac{\exp(a_j + b_j x_2) - \exp(a_j + b_j x_1)}{b_j}$$

where  $a_j$  and  $b_j$  are the intercept and slope of the  $j^{th}$  segment of  $l$ , respectively.

### **get\_f\_integral()**

This function calculates the integral under a defined interval for a provided density, given the density and a vector of points. The function returns a vector of integrals for a provided density (via the base R `integrate()` function) for each region specified by  $vector_j$  and  $vector_{j+1}$ , where  $j$  is an index for elements of a given vector.

### **is\_linear()**

This function checks if the function is linear. It generates a random sample from the domain  $D$ . If both endpoints of the inputted domain are finite, then the function simply uses those endpoints as the lower and upper limits and samples points uniformly on the domain of  $D$  to test. If either of the endpoints is infinite, then the function sets a new appropriate value as the new limit. For example, if  $D$  is unbounded on the right, we choose the new endpoint as 100 or left bound plus one, whichever is greater. Again, once we have two finite endpoints based on the domain endpoints, we sample uniformly on the interval. Then the function computes the gradient of all the sample points and checks if all of them are approximately equal.

This function is called at the beginning of `ars()` to check if the log of the given function is in a linear form.

### **sample.s()**

This function performs the actual sampling. It samples  $n$  points from  $s$  by using the inverse transform sampling method as follows. We draw a vector  $q$  of length  $n$  from the  $\text{Unif}(0, 1)$  distribution. Then, for each element of  $q$ , we find an  $x^*$  such that  $F_s(x^*) = q$ . To do so, we first have to identify the segment of  $u$  of which  $x^*$  is in the domain. Then, we calculate  $x^*$  by using the inverse CDF as follows:

$$\begin{aligned}
\int_{z_j}^{x^*} \frac{1}{I_s} \exp(u_j(t)) dt &= q - F_s(z_j) \\
\int_{z_j}^{x^*} \exp(a_j + b_j t) dt &= I_s(q - F_s(z_j)) \\
\int_{a_j + b_j z_j}^{a_j + b_j x^*} \exp(y) \frac{1}{b_j} dy &= I_s(q - F_s(z_j)) \\
[\exp(y)]_{a_j + b_j z_j}^{a_j + b_j x^*} &= b_j I_s(q - F_s(z_j)) \\
\exp(a_j + b_j x^*) - \exp(a_j + b_j z_j) &= b_j I_s(q - F_s(z_j)) \\
a_j + b_j x^* &= \log(b_j I_s(q - F_s(z_j)) + \exp(a_j + b_j z_j)) \\
x^* &= \frac{\log(b_j I_s(q - F_s(z_j)) + \exp(a_j + b_j z_j)) - a_j}{b_j}
\end{aligned}$$

Here,  $I_S$  represents the normalization factor for  $S$  and  $F_S$  is the CDF of  $S$ .

Using this formula as derived, we can convert our  $q$  vector into a vector of  $x^*$ . We return this vector of  $x^*$ s.

This function is called within `ars()` on different batch sizes to sample a vector of  $x^*$ s, which we then check for rejection or acceptance.

## ars()

This is the main function of the package. It takes in a density function `FUN` and optionally, a sample size `n`, a domain `D`, and a verbose argument which if `TRUE` will print out information on how many samples were accepted and rejected in each batch iteration.

The `ars()` function makes use of all the other functions described above to initialize `x` and then calculate `z`, `u`, `l`, and sample from `s` during each batch iteration. After getting a sample using `sample.s()`, it performs the squeezing test to ascertain which  $x^*$ s to accept and which to reject/add to the abscissae.

We check the log concavity of the provided density as `ars()` iterates over different batches (see below). We compare the integrals of the following: the provided density function, the upper hull, and the lower hull. If the density is log concave, then for all provided intervals, the upper hull will be above the density and the lower hull will be below the density. If either of these conditions is not met, then the main function will terminate.

One feature we added to `ars()` is that rather than sampling one value at a time from `sample.s()`, it samples a batch of size `batch.size` each iteration. The `batch.size` variable is initialized at 1 but is doubled each time all  $x^*$ s are accepted following the first squeezing test. This speeds up the sampling significantly.

## Tests

### Tests for Modular Components

We wrote tests for several of the modular components of our `ars()` implementation, as listed below:

- A test to check that the calculated intersections, `z`, of sequential tangent lines of concave log-density `h` are within `D`
- A test to check that the functions used to build the upper bound function outputted the correct slopes and tangent lines.
- A test to ensure that the function `get_s_integral()` outputted an integral that was larger than the integral of the density.
- A test to ensure that the function `get_l_integral()` outputs the appropriate lower bound integral.

## Kolmogorov-Smirnov Tests

To ensure the validity of samples generated from the `ars()` function, we ran a series of Kolmogorov-Smirnov tests which test for the closeness between an empirical CDF and a given distribution function in a nonparametric manner. We used the Kolmogorov-Smirnov test to test whether our `ars()` function returned a sample whose empirical CDF was close to the true CDF for a number of common log-concave distributions.

We ran the Kolmogorov-Smirnov test for the following distributions: Normal(0,1), Exponential(1), Beta(1,1), Beta(2,2), Gamma(1,1), Gamma(2,1),  $\chi^2_2$ ,  $\chi^2_3$ , and Uniform(0,1), each at a significance level of 0.05. For each test, we made sure that the p-value was greater than the significance level, suggesting that the null hypothesis, which is that the sample is drawn from the reference distribution, was true. For all tests, the p-values were greater than the default significance level.

## Team Member Contributions

Vaibhav coded functions to build U and L using the formulae provided in Gilks, calculate the integral under S using analytical solutions, and sample from S by building the inverse CDF of S and solving analytically. He also helped put together the main `ars()` function along with his teammates and especially helped with speeding up the function by enabling the sampling of points in batches. Vaibhav wrote several tests, primarily to test that the creation of the upper bound was being conducted correctly.

Colin

Brandon coded functions to calculate z (as provided by Gilks et al) and to check for density log concavity. coding of these log concavity checks was completed in parallel with Vaibhav. Brandon also included assertions within the relevant functions for troubleshooting user input.

Jonathan coded the test functions about the results of the `ars` function using the Kolmogorov-Smirnov tests with different parameters so strengthening the robustness of the code for main functions. He also created the `ars` package template and compiled them in one package and provided the descriptions for the `ars` functions.

## Package

The package is located at [www.github.com/vaibhavram/ars](https://www.github.com/vaibhavram/ars). You can install it using:

```
devtools::install_github("vaibhavram/ars")
library(ars)
```