

# Monte Carlo Simulation Tools for REDD+ Uncertainty Estimates

2024-12-19

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## 1. Introduction

When preparing for Monte Carlo simulations, it is good practice first to examine descriptive statistics of the data to characterize the empirical distributions of input variables. This preliminary analysis should include statistical tests of normality, along with visualizations of univariate distributions. Recommended visualizations include histograms, kernel density plots, and Q-Q plots. Together, these tools provide insights into the data’s shape, spread, symmetry, skewness, and potential outliers.

In particular, these visualizations help auditors to confirm levels of bias in the dataset. They also allow for quicker evaluations of how the proponent has addressed these biases in their subsequent calculations.

Incorporating these distribution assessments early in the process will likely reduce overall findings from VVB’s and registries. This is because distribution analysis is critical to selecting the appropriate functions in SimVoi, ensuring more accurate Monte Carlo estimates. In effect, bias corrections are incorporated, reducing uncertainty in the final results and improving confidence in the jurisdiction’s claims of nationwide emissions reductions.

The following guide summarizes the most commonly reported discrete and continuous distributions, their statistical properties, and possible case examples.

*Table 1: Continuous data distributions, and example use cases for Monte Carlo simulations.*

Cont. Distributions	Description of statistical criteria and common use cases
Bernoulli	Probability of a binary outcome with two possible results, such as success/failure, true/false, yes/no. E.g Probability of getting heads when flipping a single coin.
Binomial	Describes the probability of specific number of successes occurring within fixed set of independent Bernoulli trials. E.g Number of heads in 10 coin flips.

Poisson	Probability of count data, #no. of occurrences of independent events occurring within fixed time period or space. E.g #no. of customers arriving at a store per hour.
Geometric	# of failures until first success. E.g., calls until a sale.
Neg. Binomial	# of failures until r succeeds (overdispersed Poisson).
Discrete Uniform	All finite outcomes equally likely. E.g., rolling a fair die.
Normal (Gaussian)	Symmetrical “bell curve.” E.g., human heights.
Lognormal	Right-skewed; log(variable) ~ Normal. E.g., incomes.
Exponential	Time between Poisson events. E.g., arrival times.
Continuous Uniform	All values in [a,b] equally likely. E.g., random number gen.
Chi-Square	Used in hypothesis tests (e.g., goodness-of-fit).
t-Distribution	Small samples, unknown population SD.
Weibull	Reliability or lifespans.
Gamma	Models skewed data, e.g., wait times.

Table 2: Discrete data distributions, and example use cases for Monte Carlo simulations.

Discrete Distributions	Descriptions
Bernoulli	Binary outcome (success/failure). E.g., a single coin flip.
Binomial	# of successes in n Bernoulli trials. E.g., heads in 10 flips.
Poisson	# of events in a fixed interval. E.g., arrivals per hour.
Geometric	# of failures until first success. E.g., calls until a sale.

## 2. Method

### Import data

```

1 # Point this to the correct path where your file is located:
2 workbook = "./data/art/GuyanaARTWorkbookMC-thru2022-April2024_values.xlsx"
3 CarbonStocks = readxl::read_excel(workbook, "CarbonStocks") |>
4   janitor::clean_names() |>
5   mutate(across(where(is.numeric), ~round(.x, 1)))
6 CarbonStocks_MC = readxl::read_excel(workbook, "CarbonStocks (MC)") |>
7   janitor::clean_names() |>
8   mutate(across(where(is.numeric), ~round(.x, 1)))
9
10 DeforestationEF = readxl::read_excel(workbook, "Deforestation EFs") |>
11   mutate(across(where(is.numeric), ~round(.x, 1)))
12 DeforestationEF_MC = readxl::read_excel(workbook, "Deforestation EFs (MC)") |>
13   mutate(across(where(is.numeric), ~round(.x, 1)))
14
15 DegradationEF = readxl::read_excel(workbook, "Degradation EFs") |>
16   mutate(across(where(is.numeric), ~round(.x, 1)))
17 DegradationEF_MC = readxl::read_excel(workbook, "Degradation EFs (MC)") |>
18   mutate(across(where(is.numeric), ~round(.x, 1)))
19
20 ActivityData = readxl::read_excel(workbook, "Activity Data") |>
21   mutate(across(where(is.numeric), ~round(.x, 1)))
22 ActivityData_MC = readxl::read_excel(workbook, "Activity Data (MC)") |>
23   mutate(across(where(is.numeric), ~round(.x, 1)))
24
25 Emissions = readxl::read_excel(workbook, "CarbonStocks") |>

```

```

26     mutate(across(where(is.numeric), ~round(.x, 1)))
27 Emissions_MC = readxl::read_excel(workbook, "CarbonStocks (MC)") |>
28     mutate(across(where(is.numeric), ~round(.x, 1)))
29
30 # Visualize
31 flextable(head(CarbonStocks[, 1:8])) |>
32     fontsize(size = 8, part = "all")

```

x1	ag_tree_t_c_ha	bg_tree_t_c_ha	saplings_t_c_ha	standing_dead_woody_t_c_ha	dead_woody_t_c_ha	carbon_pools_t_c_ha	litter_t_c_ha
mean of all plots (calculated)	205.8	48.3	3.7	2.6	8.6	269.0	3.3
std. dev	60.4	14.3	2.0	4.0	8.1	75.2	1.3
minimum	91.6	21.2	0.5	0.0	0.0		1.2
maximum	353.7	83.1	18.8	13.7	42.3		8.7
90% CI	9.2	2.2	0.3	0.6	1.2	11.5	0.2
CI as % of mean	0.0	0.0	0.1	0.2	0.1	0.0	

```

1 flextable(head(CarbonStocks_MC[, 1:8])) |>
2     fontsize(size = 8, part = "all")

```

x1	ag_tree_t_c_ha	bg_tree_t_c_ha	saplings_t_c_ha	standing_dead_woody_t_c_ha	dead_woody_t_c_ha	carbon_pools_t_c_ha	litter_t_c_ha
tC/ha	181.1	65.0	3.5	7.3	17.1		3.7
tCO2/ha	664.2	238.2	12.8	26.9	62.6		13.7

```

1 flextable(head(CarbonStocks[, 1:8])) |>
2     fontsize(size = 8, part = "all")

```

x1	ag_tree_t_c_ha	bg_tree_t_c_ha	saplings_t_c_ha	standing_dead_woody_t_c_ha	dead_woody_t_c_ha	carbon_pools_t_c_ha	litter_t_c_ha
mean of all plots (calculated)	205.8	48.3	3.7	2.6	8.6	269.0	3.3
std. dev	60.4	14.3	2.0	4.0	8.1	75.2	1.3
minimum	91.6	21.2	0.5	0.0	0.0		1.2
maximum	353.7	83.1	18.8	13.7	42.3		8.7
90% CI	9.2	2.2	0.3	0.6	1.2	11.5	0.2
CI as % of mean	0.0	0.0	0.1	0.2	0.1	0.0	

```

1 flextable(head(CarbonStocks_MC[, 1:8])) |>
2     fontsize(size = 8, part = "all")

```

x1	ag_tree_t_c_ha	bg_tree_t_c_ha	saplings_t_c_ha	standing_dead_wood_t_c_ha	lying_dead_wood_t_c_ha	carbon_pools_litter_t_c_ha
tC/ha	181.1	65.0	3.5	7.3	17.1	3.7
tCO2/ha	664.2	238.2	12.8	26.9	62.6	13.7

```

1 # (Optionally un-comment these to view other data frames)
2 # flextable(head(DeforestationEF_MC[, 1:8])) |> fontsize(size = 8, part =
3 # 'all') flextable(head(DeforestationEF[, 1:8])) |> fontsize(size = 8, part =
4 # 'all') flextable(head(ActivityData[, 1:8])) |> fontsize(size = 8, part =
5 # 'all') flextable(head(ActivityData_MC[, 1:8])) |> fontsize(size = 8, part =
6 # 'all') flextable(head(Emissions[, 1:8])) |> fontsize(size = 8, part = 'all')
7 # flextable(head(Emissions_MC[, 1:8])) |> fontsize(size = 8, part = 'all')
8 # flextable(head(DegradationEF[, 1:8])) |> fontsize(size = 8, part = 'all')
9 # flextable(head(DegradationEF_MC[, 1:8])) |> fontsize(size = 8, part = 'all')
10 # dplyr::glimpse(CarbonStocks)

```

## Tidy data

Achieving tidy data is crucial for robust analysis, particularly when working with datasets imported from Excel, which often require adjustments to column names, strata labels, data types, and row layouts, especially when summary statistics are presented in non-standard formats. Begin by identifying the relevant rows and columns for each pool, specifically those containing mean, standard deviation, minimum, maximum, and confidence interval values. The `dplyr` package provides a powerful and efficient means for manipulating dataframes, facilitating these necessary adjustments.

Assuming the rows in the “CarbonStocks\_MC” sheet maintain a consistent order, operations can be performed by simply referencing the dataframe name. A common approach involves reshaping the data so that each row represents a “Statistic,” such as mean or standard deviation, and each column corresponds to a carbon pool, like “AG Tree” or “BG Tree.”

Initially, select the columns pertinent to your carbon pools, for instance, those named “AG Tree (tC/ha)” or “BG Tree (tC/ha),” and rename them to align with the “SimVoi” workbook. Subsequently, extract the rows containing the summary statistics, typically the first few rows, and proceed to reshape the data. Note that direct renaming of row values, such as assigning “mean” to the second row, must be explicitly performed if required, as it is not automatically handled.

Throughout this process, thorough inspection of the data using `view(CarbonStocks)` or `glimpse(CarbonStocks)` after each operation is strongly recommended. This ensures accurate mapping of rows and columns to the “Statistic” labels.

To effectively transpose the data and transition between wide and long formats, utilize the `tidyr` package’s `pivot_longer()` and `pivot_wider()` functions. These functions facilitate the transformation of data such that each row contains a data value, such as mean, standard deviation, or minimum, and each column represents a variable or carbon pool. Be mindful of potential missing or incorrect column names, often indicated by placeholder names like “...1” generated by `readxl::read_excel()`; these must be renamed. Finally, pivot the data from long format back to wide, ensuring that “Statistic” becomes a distinct column and the carbon pools, such as “AG\_Tree” and “BG\_Tree,” are represented as separate variable columns.

```
# CarbonStocks <- CarbonStocks |> rename(Statistic = `...1`)
```

```
CarbonStocks <- CarbonStocks %>%
  select(Statistic = 1, AG_Tree = 2, BG_Tree = 3, Saplings = 4, StandingDeadWood = 5,
         LyingDeadWood = 6, SumCarbonNoLitter = 7, Litter = 8, SumCpoolWlitter = 9,
         SumCO2e = 10, Soil_tC_ha = 11, SumALL_POOLS_CO2eha = 12, SumABGBLiveTree = 13) %>%
  slice(1:9)
# Rename missing column names
```

```

# Convert from wide to long: 'Statistic' will define the row identity
CarbonStocks_long <- CarbonStocks |>
  tidyr::pivot_longer(cols = -Statistic, names_to = "Pool", values_to = "Value") |>
  mutate(Value = as.numeric(Value))

# Convert from long back to wide format:
CarbonStocks_wide <- CarbonStocks_long %>%
  pivot_wider(names_from = Statistic, values_from = Value)

# Inspect the final structure
CarbonStocks_wide

# Example summarizing a particular column:
CarbonStocks_wide %>%
  summarise(Mean_AGTree = mean(`AG Tree (tC/ha)`, na.rm = TRUE), SD_AGTree = sd(`AG Tree (tC/ha)`,
    na.rm = TRUE))

# Transpose to long dataframe: flipping rows w/ columns
CarbonStocks_long <- CarbonStocks |>
  tidyr::pivot_longer(cols = -Statistic, names_to = "Pool", values_to = "Value") |>
  mutate(Value = as.numeric(Value))

# Pivot back to wide dataframe & "Statistic" becomes a row:
CarbonStocks_wide <- CarbonStocks_long %>%
  pivot_wider(names_from = Statistic, values_from = Value)
# Inspect
CarbonStocks_wide

CarbonStocks_wide |>
  summarise(Mean_AGTree = mean(`AG Tree (tC/ha)`, na.rm = TRUE), SD_AGTree = sd(`AG Tree (tC/ha)`,
    na.rm = TRUE))

```

## Descriptive Statistics

### Shapiro-Wilk test for normality

```
shapiro.test(CarbonStocks$AG Tree (tC/ha))
```

## Distribution Analysis

### Example approximate histogram

```
hist(AG_draws, breaks=30, main="AG Tree (approx. distribution)") qqnorm(AG_draws) qq-
line(AG_draws, col="red")
```

The Coefficient of Variation CV is a standardized, unit-less measure of dispersion defined as the ratio of the standard deviation to the mean, typically expressed as a percentage. This standardization allows for meaningful comparisons of variability across datasets or scales, regardless of the underlying units, offering helpful tool for assessing novel data from periodic field inventories or mapping updates.

$$CV = \frac{\sigma}{\mu} \times 100\%$$

$$CV_{\%} = 100 \times \frac{\text{std. dev}}{\text{mean of all plots (calculated)}}$$

For these carbon stocks, a higher CV indicates greater relative variability or “scatter” in the data. While the CV is a useful indicator of dispersion and can signal potential non-normality, it does not provide any information on the direction of skew in the distribution.

In our analysis, the CV was computed below within a helper function called `calc_derived_stats`. This function not only calculates the CV but also compares the reported 90% confidence interval with the standard deviation, which, under normality, should approximate to  $\pm 1.645 \times \text{SD}$ . This iterative scoring helps assess the internal consistency of the reported descriptive statistics.

*# Helper function of derived descriptive statistics:*

```
calc_derived_stats <- function(df) {
  df %>%
    mutate(
      CV_percent = 100 * (`std. dev` / `mean of all plots (calculated)`),
      sd_implied_by_90CI = `90% CI` / 1.645,
      SDs_below_mean = (`mean of all plots (calculated)` - minimum) / `std. dev`,
      SDs_above_mean = (maximum - `mean of all plots (calculated)`) / `std. dev`
    )
}
```

```
CarbonStocks_stats <- calc_derived_stats(CarbonStocks_wide)
```

*#CarbonStocks\_stats # Remember to inspect new variables*

*# Custom function to simulate from each row (assuming truncnormal)*

```
simulate_truncnorm_from_summary <- function(
  mean_val, sd_val, min_val=0, max_val=Inf,
  # We loaded here the 'truncnorm' package from main cran libraries, I will add to in-line
  n_draws=10000) {
  draws <- truncnorm::rtruncnorm(
    n      = n_draws,
    a      = min_val,
    b      = max_val,
    mean   = mean_val,
    sd     = sd_val
  )
  # Return vector of draws
  return(draws)
}
```

*# Repeat for AG\_Tree*

```
ag_tree_stats <- CarbonStocks_stats %>% filter(Pool == "AG_Tree")
AG_mean <- ag_tree_stats$`mean of all plots (calculated)`
AG_sd   <- ag_tree_stats$`std. dev`
AG_min  <- ag_tree_stats$minimum
AG_max  <- ag_tree_stats$maximum
```

*# We may vote to do a = 0 if we never allow negative carbon:*

```
AG_draws <- simulate_truncnorm_from_summary(
  mean_val = AG_mean,
  sd_val   = AG_sd,
  min_val  = 0,      # or AG_min if you prefer
```

```

    max_val = Inf,
    n_draws = 10000
  )

  # Compare results:
  mean(AG_draws)
  sd(AG_draws)
  min(AG_draws)
  max(AG_draws)
  quantile(AG_draws, probs = c(0.05, 0.95))

  # Quick histogram of the draws
  hist(AG_draws, breaks=40, col="skyblue",
       main="Truncated Normal draws for AG Tree",
       xlab="AG Tree (tC/ha)")

  # If you want to do this for each carbon pool in a loop,
  # you can add a small function:

  simulate_all_pools <- function(df, n_draws=10000) {
    # df is your cs_stats data frame
    # Return a named list of random draws
    out <- list()
    for (i in seq_len(nrow(df))) {
      rowi <- df[i, ]
      pool_name <- rowi$Pool
      mean_val <- rowi$`mean of all plots (calculated)`
      sd_val <- rowi$`std. dev`
      # Use zero for min bound; or rowi$minimum if you want to
      # replicate the workbook min
      draws <- rtruncnorm(
        n=n_draws,
        a=0,
        b=Inf,
        mean=mean_val,
        sd=sd_val
      )
      out[[pool_name]] <- draws
    }
    return(out)
  }

  all_draws <- simulate_all_pools(CarbonStocks_st_stats, n_draws=10000)

  names(l_draws)

  ggplot(data.frame(AG_draws), aes(x = AG_draws)) +
    geom_histogram(aes(y = ..density..), bins = 50, fill = "skyblue", alpha = 0.7) +
    geom_density(col = "red") +

```

```
labs(title = "Monte Carlo Simulation of AG Tree Carbon Pool",
     x = "Carbon Stock (tC/ha)", y = "Density")
```

## Replicating SimVoi

We utilize the replicate function to repeat a simulation following a randomized normally truncated multiple times with replicate(n=10000, while determining the size of the sampled subset with rnorm(n=100. The first model explores sample size parameters only, replication parameters are tested below this in comparisons.

```
MEAN = CarbonStocks$`AG Tree (tC/ha)`[1]
SD = CarbonStocks$`AG Tree (tC/ha)`[2]

randtruncnormal_sim_10000 <- rnorm(n = 10000, mean = MEAN, sd = SD)
hist(randtruncnormal_sim_10000, freq = F)
AG_Tree_tC_ha = mean(randtruncnormal_sim_10000)
AG_Tree_tCO2_ha = AG_Tree_tC_ha * (44/12)
AG_Tree_tC_ha
AG_Tree_tCO2_ha
# curve(dnorm(x, mean=MEAN, sd=SD), from=0, to=450, add=T, col='red')
```

## Compare simulations

```
# 10,000 simulations sampling 10 observations
randtruncnormal_sim_10000_10 = replicate(n = 10000, rnorm(n = 10, mean = MEAN, sd = SD))
hist(apply(X = randtruncnormal_sim_10000_10, MARGIN = 2, FUN = mean))
sd(apply(X = randtruncnormal_sim_10000_10, MARGIN = 2, FUN = mean))
mean(apply(X = randtruncnormal_sim_10000_10, MARGIN = 2, FUN = mean))
(mean(apply(X = randtruncnormal_sim_10000_10, MARGIN = 2, FUN = mean))) * (44/12)

# 10,000 simulations sampling 100 observations
randtruncnormal_sim_10000_100 = replicate(n = 10000, rnorm(n = 100, mean = MEAN,
  sd = SD))
hist(apply(X = randtruncnormal_sim_10000_100, MARGIN = 2, FUN = mean))
sd(apply(X = randtruncnormal_sim_10000_100, MARGIN = 2, FUN = mean))
mean(apply(X = randtruncnormal_sim_10000_100, MARGIN = 2, FUN = mean))
(mean(apply(X = randtruncnormal_sim_10000_100, MARGIN = 2, FUN = mean))) * (44/12)

# 10,000 simulations sampling 1,000 observations
randtruncnormal_sim_10000_1000 = replicate(n = 10000, rnorm(n = 1000, mean = MEAN,
  sd = SD))
hist(apply(X = randtruncnormal_sim_10000_1000, MARGIN = 2, FUN = mean))
sd(apply(X = randtruncnormal_sim_10000_1000, MARGIN = 2, FUN = mean))
mean(apply(X = randtruncnormal_sim_10000_1000, MARGIN = 2, FUN = mean))
(mean(apply(X = randtruncnormal_sim_10000_1000, MARGIN = 2, FUN = mean))) * (44/12)

# 10,000 simulations sampling 10,000 observations
randtruncnormal_sim_10000_10000 = replicate(n = 10000, rnorm(n = 10000, mean = MEAN,
  sd = SD))
hist(apply(X = randtruncnormal_sim_10000_10000, MARGIN = 2, FUN = mean))
sd(apply(X = randtruncnormal_sim_10000_10000, MARGIN = 2, FUN = mean))
mean(apply(X = randtruncnormal_sim_10000_10000, MARGIN = 2, FUN = mean))
(mean(apply(X = randtruncnormal_sim_10000_10000, MARGIN = 2, FUN = mean))) * (44/12)

devtools::session_info()

- Session info -----
```



```

setting  value
version  R version 4.3.0 (2023-04-21)
os       macOS 15.3.2
system   aarch64, darwin20
ui       X11
language (EN)
collate  en_US.UTF-8
ctype    en_US.UTF-8
tz       America/Vancouver
date     2025-03-16
pandoc   3.6.1 @ /usr/local/bin/ (via rmarkdown)

```

```

- Packages -----
package      * version      date (UTC) lib source
abind         1.4-8        2024-09-12 [1] CRAN (R 4.3.3)
animation    * 2.7         2021-10-07 [1] CRAN (R 4.3.3)
askpass       1.2.1        2024-10-04 [1] CRAN (R 4.3.3)
assertthat    0.2.1        2019-03-21 [1] CRAN (R 4.3.0)
backports     1.5.0        2024-05-23 [1] CRAN (R 4.3.3)
BIOMASS       * 2.2.3       2025-02-24 [1] CRAN (R 4.3.3)
boot          1.3-31       2024-08-28 [1] CRAN (R 4.3.3)
broom         * 1.0.7       2024-09-26 [1] CRAN (R 4.3.3)
c2z           * 0.2.0       2023-08-10 [1] CRAN (R 4.3.0)
cachem        1.1.0        2024-05-16 [1] CRAN (R 4.3.3)
car           3.1-3        2024-09-27 [1] CRAN (R 4.3.3)
carData       3.0-5        2022-01-06 [1] CRAN (R 4.3.0)
caret         * 7.0-1       2024-12-10 [1] CRAN (R 4.3.3)
cellranger    1.1.0        2016-07-27 [1] CRAN (R 4.3.0)
chromote      0.4.0        2025-01-25 [1] CRAN (R 4.3.3)
class         7.3-23       2025-01-01 [1] CRAN (R 4.3.3)
classInt      0.4-11       2025-01-08 [1] CRAN (R 4.3.3)
cli           3.6.3        2024-06-21 [1] CRAN (R 4.3.3)
codetools     0.2-20       2024-03-31 [1] CRAN (R 4.3.1)
colorspace    2.1-1        2024-07-26 [1] CRAN (R 4.3.3)
data.table    1.16.4       2024-12-06 [1] CRAN (R 4.3.3)
dataMaid      * 1.4.1       2021-10-08 [1] CRAN (R 4.3.0)
DBI           1.2.3        2024-06-02 [1] CRAN (R 4.3.3)
DEoptimR      1.1-3-1      2024-11-23 [1] CRAN (R 4.3.3)
DescTools     * 0.99.59     2025-01-26 [1] CRAN (R 4.3.3)
devtools      2.4.5        2022-10-11 [1] CRAN (R 4.3.0)
dials         * 1.3.0       2024-07-30 [1] CRAN (R 4.3.3)
DiceDesign    1.10         2023-12-07 [1] CRAN (R 4.3.1)
digest        0.6.37       2024-08-19 [1] CRAN (R 4.3.3)
dplyr         * 1.1.4       2023-11-17 [1] CRAN (R 4.3.1)
e1071         1.7-16       2024-09-16 [1] CRAN (R 4.3.3)
easypackages  0.1.0        2016-12-05 [1] CRAN (R 4.3.0)
ellipsis      0.3.2        2021-04-29 [1] CRAN (R 4.3.0)
evaluate      1.0.3        2025-01-10 [1] CRAN (R 4.3.3)
Exact         3.3          2024-07-21 [1] CRAN (R 4.3.3)
expm          1.0-0        2024-08-19 [1] CRAN (R 4.3.3)
extrafont     * 0.19        2023-01-18 [1] CRAN (R 4.3.3)
extrafontdb   1.0          2012-06-11 [1] CRAN (R 4.3.3)
fastmap       1.2.0        2024-05-15 [1] CRAN (R 4.3.3)
flextable     * 0.9.7       2024-10-27 [1] CRAN (R 4.3.3)

```

fontBitstreamVera	0.1.1	2017-02-01	[1]	CRAN	(R 4.3.3)
fontLiberation	0.1.0	2016-10-15	[1]	CRAN	(R 4.3.3)
fontquiver	0.2.1	2017-02-01	[1]	CRAN	(R 4.3.3)
forcats	* 1.0.0	2023-01-29	[1]	CRAN	(R 4.3.0)
foreach	1.5.2	2022-02-02	[1]	CRAN	(R 4.3.0)
formatR	* 1.14	2023-01-17	[1]	CRAN	(R 4.3.3)
Formula	1.2-5	2023-02-24	[1]	CRAN	(R 4.3.0)
fs	1.6.5	2024-10-30	[1]	CRAN	(R 4.3.3)
furrr	0.3.1	2022-08-15	[1]	CRAN	(R 4.3.0)
future	1.34.0	2024-07-29	[1]	CRAN	(R 4.3.3)
future.apply	1.11.3	2024-10-27	[1]	CRAN	(R 4.3.3)
gdtools	0.4.1	2024-11-04	[1]	CRAN	(R 4.3.3)
generics	0.1.3	2022-07-05	[1]	CRAN	(R 4.3.0)
ggplot2	* 3.5.1	2024-04-23	[1]	CRAN	(R 4.3.1)
gld	2.6.7	2025-01-17	[1]	CRAN	(R 4.3.3)
globals	0.16.3	2024-03-08	[1]	CRAN	(R 4.3.1)
glue	1.8.0	2024-09-30	[1]	CRAN	(R 4.3.3)
goftest	1.2-3	2021-10-07	[1]	CRAN	(R 4.3.3)
gower	1.0.2	2024-12-17	[1]	CRAN	(R 4.3.3)
GPfit	1.0-8	2019-02-08	[1]	CRAN	(R 4.3.0)
gridExtra	2.3	2017-09-09	[1]	CRAN	(R 4.3.0)
gtable	0.3.6	2024-10-25	[1]	CRAN	(R 4.3.3)
hardhat	1.4.0	2024-06-02	[1]	CRAN	(R 4.3.3)
haven	2.5.4	2023-11-30	[1]	CRAN	(R 4.3.1)
hms	1.1.3	2023-03-21	[1]	CRAN	(R 4.3.0)
htmltools	* 0.5.8.1	2024-04-04	[1]	CRAN	(R 4.3.1)
htmlwidgets	1.6.4	2023-12-06	[1]	CRAN	(R 4.3.1)
httpuv	1.6.15	2024-03-26	[1]	CRAN	(R 4.3.1)
httr	1.4.7	2023-08-15	[1]	CRAN	(R 4.3.0)
infer	* 1.0.7	2024-03-25	[1]	CRAN	(R 4.3.1)
ipred	0.9-15	2024-07-18	[1]	CRAN	(R 4.3.3)
iterators	1.0.14	2022-02-05	[1]	CRAN	(R 4.3.0)
janitor	* 2.2.1	2024-12-22	[1]	CRAN	(R 4.3.3)
jsonlite	* 1.8.9	2024-09-20	[1]	CRAN	(R 4.3.3)
kableExtra	* 1.4.0	2024-01-24	[1]	CRAN	(R 4.3.1)
kernlab	* 0.9-33	2024-08-13	[1]	CRAN	(R 4.3.3)
KernSmooth	2.23-26	2025-01-01	[1]	CRAN	(R 4.3.3)
knitr	* 1.49	2024-11-08	[1]	CRAN	(R 4.3.3)
later	1.4.1	2024-11-27	[1]	CRAN	(R 4.3.3)
lattice	* 0.22-6	2024-03-20	[1]	CRAN	(R 4.3.1)
lava	1.8.1	2025-01-12	[1]	CRAN	(R 4.3.3)
lazyeval	0.2.2	2019-03-15	[1]	CRAN	(R 4.3.0)
lhs	1.2.0	2024-06-30	[1]	CRAN	(R 4.3.3)
lifecycle	1.0.4	2023-11-07	[1]	CRAN	(R 4.3.1)
listenv	0.9.1	2024-01-29	[1]	CRAN	(R 4.3.1)
lmom	3.2	2024-09-30	[1]	CRAN	(R 4.3.3)
lubridate	* 1.9.4	2024-12-08	[1]	CRAN	(R 4.3.3)
magrittr	2.0.3	2022-03-30	[1]	CRAN	(R 4.3.0)
MASS	7.3-58.4	2023-03-07	[2]	CRAN	(R 4.3.0)
Matrix	1.6-5	2024-01-11	[1]	CRAN	(R 4.3.1)
memoise	2.0.1	2021-11-26	[1]	CRAN	(R 4.3.0)
mime	0.12	2021-09-28	[1]	CRAN	(R 4.3.0)
miniUI	0.1.1.1	2018-05-18	[1]	CRAN	(R 4.3.0)
minpack.lm	1.2-4	2023-09-11	[1]	CRAN	(R 4.3.3)

mnormt	2.1.1	2022-09-26	[1]	CRAN	(R 4.3.0)
modeldata	* 1.4.0	2024-06-19	[1]	CRAN	(R 4.3.3)
ModelMetrics	1.2.2.2	2020-03-17	[1]	CRAN	(R 4.3.0)
munsell	0.5.1	2024-04-01	[1]	CRAN	(R 4.3.1)
mvtnorm	1.3-3	2025-01-10	[1]	CRAN	(R 4.3.3)
nlme	3.1-166	2024-08-14	[1]	CRAN	(R 4.3.3)
nnet	7.3-20	2025-01-01	[1]	CRAN	(R 4.3.3)
nortest	1.0-4	2015-07-30	[1]	CRAN	(R 4.3.3)
officer	0.6.7	2024-10-09	[1]	CRAN	(R 4.3.3)
olsrr	* 0.6.1	2024-11-06	[1]	CRAN	(R 4.3.3)
openssl	2.3.1	2025-01-09	[1]	CRAN	(R 4.3.3)
pander	0.6.6	2025-03-01	[1]	CRAN	(R 4.3.3)
parallelly	1.41.0	2024-12-18	[1]	CRAN	(R 4.3.3)
parsnip	* 1.2.1	2024-03-22	[1]	CRAN	(R 4.3.1)
pillar	1.10.1	2025-01-07	[1]	CRAN	(R 4.3.3)
pkgbuild	1.4.6	2025-01-16	[1]	CRAN	(R 4.3.3)
pkgconfig	2.0.3	2019-09-22	[1]	CRAN	(R 4.3.0)
pkgload	1.4.0	2024-06-28	[1]	CRAN	(R 4.3.3)
plotly	* 4.10.4	2024-01-13	[1]	CRAN	(R 4.3.1)
plyr	1.8.9	2023-10-02	[1]	CRAN	(R 4.3.1)
pROC	1.18.5	2023-11-01	[1]	CRAN	(R 4.3.1)
processx	3.8.5	2025-01-08	[1]	CRAN	(R 4.3.3)
prodlim	2024.06.25	2024-06-24	[1]	CRAN	(R 4.3.3)
profvis	0.4.0	2024-09-20	[1]	CRAN	(R 4.3.3)
promises	1.3.2	2024-11-28	[1]	CRAN	(R 4.3.3)
proxy	0.4-27	2022-06-09	[1]	CRAN	(R 4.3.0)
ps	1.8.1	2024-10-28	[1]	CRAN	(R 4.3.3)
psych	* 2.4.12	2024-12-23	[1]	CRAN	(R 4.3.3)
purrr	* 1.0.2	2023-08-10	[1]	CRAN	(R 4.3.0)
R6	2.5.1	2021-08-19	[1]	CRAN	(R 4.3.0)
ragg	1.3.3	2024-09-11	[1]	CRAN	(R 4.3.3)
rappdirs	0.3.3	2021-01-31	[1]	CRAN	(R 4.3.0)
RColorBrewer	* 1.1-3	2022-04-03	[1]	CRAN	(R 4.3.0)
Rcpp	1.0.14	2025-01-12	[1]	CRAN	(R 4.3.3)
readr	* 2.1.5	2024-01-10	[1]	CRAN	(R 4.3.1)
readxl	* 1.4.3	2023-07-06	[1]	CRAN	(R 4.3.0)
recipes	* 1.1.0	2024-07-04	[1]	CRAN	(R 4.3.3)
remotes	2.5.0	2024-03-17	[1]	CRAN	(R 4.3.1)
reshape2	1.4.4	2020-04-09	[1]	CRAN	(R 4.3.0)
rlang	1.1.4	2024-06-04	[1]	CRAN	(R 4.3.3)
rmarkdown	* 2.29	2024-11-04	[1]	CRAN	(R 4.3.3)
robustbase	0.99-4-1	2024-09-27	[1]	CRAN	(R 4.3.3)
rootSolve	1.8.2.4	2023-09-21	[1]	CRAN	(R 4.3.3)
rpart	4.1.24	2025-01-07	[1]	CRAN	(R 4.3.3)
rsample	* 1.2.1	2024-03-25	[1]	CRAN	(R 4.3.1)
rstudioapi	0.17.1	2024-10-22	[1]	CRAN	(R 4.3.3)
Rttf2pt1	1.3.12	2023-01-22	[1]	CRAN	(R 4.3.3)
rvest	1.0.4	2024-02-12	[1]	CRAN	(R 4.3.1)
scales	* 1.3.0	2023-11-28	[1]	CRAN	(R 4.3.1)
sessioninfo	1.2.2	2021-12-06	[1]	CRAN	(R 4.3.0)
sf	1.0-19	2024-11-05	[1]	CRAN	(R 4.3.3)
shiny	1.10.0	2024-12-14	[1]	CRAN	(R 4.3.3)
snakecase	0.11.1	2023-08-27	[1]	CRAN	(R 4.3.0)
stringi	1.8.4	2024-05-06	[1]	CRAN	(R 4.3.1)

stringr	* 1.5.1	2023-11-14	[1]	CRAN	(R 4.3.1)
survival	3.8-3	2024-12-17	[1]	CRAN	(R 4.3.3)
svglite	2.1.3	2023-12-08	[1]	CRAN	(R 4.3.1)
systemfonts	1.1.0	2024-05-15	[1]	CRAN	(R 4.3.3)
terra	1.8-29	2025-02-26	[1]	CRAN	(R 4.3.3)
textshaping	0.4.1	2024-12-06	[1]	CRAN	(R 4.3.3)
tibble	* 3.2.1	2023-03-20	[1]	CRAN	(R 4.3.0)
tidymodels	* 1.2.0	2024-03-25	[1]	CRAN	(R 4.3.1)
tidyr	* 1.3.1	2024-01-24	[1]	CRAN	(R 4.3.1)
tidyselect	1.2.1	2024-03-11	[1]	CRAN	(R 4.3.1)
tidyverse	* 2.0.0	2023-02-22	[1]	CRAN	(R 4.3.0)
timechange	0.3.0	2024-01-18	[1]	CRAN	(R 4.3.1)
timeDate	4041.110	2024-09-22	[1]	CRAN	(R 4.3.3)
tinytex	* 0.54	2024-11-01	[1]	CRAN	(R 4.3.3)
truncnorm	* 1.0-9	2023-03-20	[1]	CRAN	(R 4.3.3)
tune	* 1.2.1	2024-04-18	[1]	CRAN	(R 4.3.1)
tzdb	0.4.0	2023-05-12	[1]	CRAN	(R 4.3.0)
units	0.8-5	2023-11-28	[1]	CRAN	(R 4.3.1)
urlchecker	1.0.1	2021-11-30	[1]	CRAN	(R 4.3.0)
useful	* 1.2.6.1	2023-10-24	[1]	CRAN	(R 4.3.1)
usethis	3.1.0	2024-11-26	[1]	CRAN	(R 4.3.3)
uuid	1.2-1	2024-07-29	[1]	CRAN	(R 4.3.3)
vctrs	0.6.5	2023-12-01	[1]	CRAN	(R 4.3.1)
viridisLite	0.4.2	2023-05-02	[1]	CRAN	(R 4.3.0)
webshot	* 0.5.5	2023-06-26	[1]	CRAN	(R 4.3.0)
webshot2	* 0.1.1	2023-08-11	[1]	CRAN	(R 4.3.0)
websocket	1.4.2	2024-07-22	[1]	CRAN	(R 4.3.3)
withr	3.0.2	2024-10-28	[1]	CRAN	(R 4.3.3)
workflows	* 1.1.4	2024-02-19	[1]	CRAN	(R 4.3.1)
workflowsets	* 1.1.0	2024-03-21	[1]	CRAN	(R 4.3.1)
xfun	0.50	2025-01-07	[1]	CRAN	(R 4.3.3)
xml2	1.3.6	2023-12-04	[1]	CRAN	(R 4.3.1)
xtable	1.8-4	2019-04-21	[1]	CRAN	(R 4.3.0)
yaml	2.3.10	2024-07-26	[1]	CRAN	(R 4.3.3)
yardstick	* 1.3.1	2024-03-21	[1]	CRAN	(R 4.3.1)
zip	2.3.1	2024-01-27	[1]	CRAN	(R 4.3.1)

[1] /Library/Frameworks/R.framework/Versions/4.1-arm64/Resources/library

[2] /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library

---

```
# Sys.getenv() .libPaths()
```

## References

### Annex I: SimVoi Functions & Syntax

SimVoi adds seventeen random number generator functions defined with the following syntax:

- `RandBeta(alpha,beta,,[MinValue],[MaxValue])`
- `RandBinomial(trials,probability_s)`
- `RandBiVarNormal(mean1,stdev1,mean2,stdev2,correl12)`
- `RandCumulative(value_cumulative_table)`
- `RandDiscrete(value_discrete_table)`

- `RandExponential(lambda)`
- `RandInteger(bottom,top)`
- `RandLogNormal(Mean,StDev)`
- `RandNormal(mean,standard_dev)`
- `RandPoisson(mean)`
- `RandSample(population)`
- `RandTriangular(minimum,most_likely,maximum)`
- `RandTriBeta(minimum,most_likely,maximum,[shape])`
- `RandTruncBiVarNormal(mean1,stdev1,mean2,stdev2,correl12,[min1],[max1],[min2],[max2])`
- `RandTruncLogNormal(Mean,StDev,[MinValue],[MaxValue])`
- `RandTruncNormal(Mean,StDev,[MinValue],[MaxValue])`
- `RandUniform(minimum,maximum)`

In the following, we attempt to match the SimVoi Excel formula of

```
= [1] !randtruncnormal(CarbonStocks.B2,CarbonStocks.B3,0)
```

function, as closely as random seeding allows. According to package documentation, the `RandTruncNormal()` function “Returns a random value from a truncated normal probability density function. This function can model an uncertain quantity with a bell-shaped density function where extreme values in the tails of the distribution are not desired.”

In terms of simulation parameters, “*RandTruncNormal(Mean,StDev,MinValue,MaxValue)*” uses values of *RandNormal* until a value is found between *MinValue* and *MaxValue* or until it has made 10,000 attempts.” The above formula provides a minimum value of 0, passing to the default number of simulations of 10,000.