Review of SimVoi's Monte Carlo Simulations of Guyana's REDD+ Uncertainty Esimates

Winrock International*

2024-03-24

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Objective

Using an R-based approach, this analysis replicates the Monte Carlo simulations originally performed with the SimVoi add-in in Excel.¹. It details the code used in the analysis, compares simulation results between R and Excel, and proposes next steps for enhancement based on statistical tests. Such updates are targeted to provide an effective basis for reducing uncertainty and increasing revenue from emissions credits.

SimVoi syntax

SimVoi provides seventeen random number generator functions² that are operated 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)

^{*}seamus.murphy@winrock.org, pedro.piffer@winrock.org, meyru.bhanti@winrock.org

¹Mac Add-in: https://treeplan.com/wp-content/uploads/How-To-Install-Mac-Excel-Addin.pdf

Windows Add-in: https://treeplan.com/wp-content/uploads/How-To-Install-Windows-Excel-Addin.pdf

 $^{^2 \}mathrm{SimVoi}\text{-}313\text{-}\mathrm{Guide.pdf}$

- 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 this workflow, we attempt to replicate the following SimVoi function as identified in Guyana's emissions workbook:

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

According to the documentation, RandTruncNormal() returns a random value from a truncated normal distribution, modeling an uncertain quantity with a bell-shaped density while excluding extreme tail values. If no simulation count is provided, the function repeatedly samples until a value falls between the specified minimum and maximum, or until it reaches 10,000 attempts. In the above example, only a minimum value of 0 is provided, so the default iteration limit is used.

$Import\ data$

```
workbook = "./data/art/GuyanaARTWorkbookMC-thru2022-April2024_values_V2.xlsx"

CarbonStocks = readxl::read_excel(workbook, "CarbonStocks") |>
    janitor::clean_names() |> mutate(across(where(is.numeric), ~ round(.x, 1)))

DegradationEF = readxl::read_excel(workbook, "Degradation EFs") |> janitor::clean_names()|>
    mutate(across(where(is.numeric), ~ round(.x, 1)))

ActivityData = readxl::read_excel(workbook, "Activity Data") |> janitor::clean_names() |>
    mutate(across(where(is.numeric), ~ round(.x, 1)))

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

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

flextable(ActivityData[, 1:8])|>fontsize(size=8,part="all")
```

 $Table\ 1:\ Input\ values\ from\ Carbon Stocks\ tab sheet$

Statistic	AG Tree	BG Tree	Saplings	Standing Dead	Lying Dead	Litter	Sum w/o Soil	Soil
mean	205.8	48.3	3.7	2.6	8.6	3.3	272.3	58.7
std_dev	60.4	14.3	2.0	4.0	8.1	1.3	90.0	61.5
minimum	91.6	21.2	0.5	0.0	0.0	1.2	114.4	10.1
maximum	353.7	83.1	18.8	13.7	42.3	8.7	520.3	502.4
90%_CI	9.2	2.2	0.3	0.6	1.2	0.2	N/A	11.0
CI_%_of_mean	0.0	0.0	0.1	0.2	0.1	N/A	N/A	0.2

 ${\it Table~2:~Input~values~from~Degradation~EFs~tabsheet}$

Statistic	LDF	Wood Density	LIF	For. Infrastr.	Mining	Mining Infrastr.	Infrast.
Factor_tC	1.05	0.4	46.87	NA	NA	NA	NA
StdDev_tC	0.68	0.03	8.08	NA	NA	NA	NA
CI90_tC	0.08	0	1.6	NA	NA	NA	NA
Factor_tCO2	3.85	1.47	171.84	NA	NA	NA	NA
StdDev_tCO2	2.4900000000000002	0.11	29.63	NA	NA	NA	NA
Cl90_tCO2	0.289999999999998	0.01	5.87	NA	NA	NA	NA

Statistic	LDF	Wood Density	LIF	For. Infrastr.	Mining	Mining Infrastr.	Infrast.
EF_tCO2_AD	NA	NA	NA	8.1	8.1	8.1	8.1
MAD_tCO2	NA	NA	NA	8.1	8.1	8.1	8.1

Table 3: Absolute input values from Activity Data tabsheet

drivers	units	x2011	x2012	x2013	x2014	x2015	x2016	x2017	x2018	x2019	x2020	x2021	x2022	x2023
Forestry infrastructure	ha	186	240	330	204	313	313	227	356	226	195	228	155.6	339
Agriculture	ha	41	440	424	817	379	379	477	512	246	489	216	281.6	475
Mining (medium and large scale)	ha	7340	13664	11518	10434	6782	6782	7442	7624	5821	6,452	6,825	5,264.3	5,853
Mining infrastructure	ha													
Infrastructure	ha	118	127	342	141	217	217	195	67	52	102	117	110.6	541
Settlements	ha	-	-	23	71	8	8	7	7	22	60	105	169.4	201

SimVoi replication

Please note that the number of iterations per simulation were reduced in the following tests specifically to explore treatment of convergence in SimVoi defaults operations.

$CarbonStocks\ data$

```
A_MEAN = CarbonStocks$ag_tree_t_c_ha[1]
         = CarbonStocks$ag tree t c ha[2]
   B_MEAN = CarbonStocks$bg_tree_t_c_ha[1]
        = CarbonStocks$bg_tree_t_c_ha[2]
   C_MEAN = CarbonStocks$saplings_t_c_ha[1]
        = CarbonStocks$saplings_t_c_ha[2]
   D_MEAN = CarbonStocks$standing_dead_wood_t_c_ha[1]
         = CarbonStocks$standing_dead_wood_t_c_ha[2]
   E_MEAN = CarbonStocks$lying_dead_wood_t_c_ha[1]
         = CarbonStocks$lying_dead_wood_t_c_ha[2]
10
   F_MEAN = CarbonStocks$litter_t_c_ha[1]
   F_SD
         = CarbonStocks$litter_t_c_ha[2]
   G_MEAN = CarbonStocks$sum_pools_w_o_soil[1]
        = CarbonStocks$sum pools w o soil[2]
   H_MEAN = CarbonStocks$soil_t_c_ha[1]
         = CarbonStocks$soil_t_c_ha[2]
16
   # 100 simulations sampling 1 observation each iteration.
18
   A_rtruncnormal_100 = truncnorm::rtruncnorm(n=100,91.6,353.7,A_MEAN, A_SD)
   B_rtruncnormal_100 = truncnorm::rtruncnorm(n=100,21.2,83.1,B_MEAN, B_SD)
20
   C_rtruncnormal_100 = truncnorm::rtruncnorm(n=100,0.5,18.8,C_MEAN, C_SD)
   D_rtruncnormal_100 = truncnorm::rtruncnorm(n=100,0.0,13.7,D_MEAN, D_SD)
22
   E_rtruncnormal_100 = truncnorm::rtruncnorm(n=100,0.0,42.3,E_MEAN, E_SD)
   F_rtruncnormal_100 = truncnorm::rtruncnorm(n=100,1.2,8.7,F_MEAN,F_SD)
24
   G_rtruncnormal_100 = truncnorm::rtruncnorm(n=100,114.4,520.3,G_MEAN, G_SD)
   H_rtruncnormal_100 = truncnorm::rtruncnorm(n=100,10.1,502.4,H_MEAN, H_SD)
26
27
   # --- Simulation Estimates ---
```

```
AG_tree_tC_ha
                             = mean(A rtruncnormal 100)
                             = mean(A rtruncnormal 100)*(44/12)
   AG_tree_tCO2_ha
30
   BG tree tC ha
                             = mean(B rtruncnormal 100)
31
                             = mean(B_rtruncnormal_100)*(44/12)
   BG_tree_tCO2_ha
   Saplings_tC_ha
                             = mean(C rtruncnormal 100)
33
   Saplings tCO2 ha
                             = mean(C rtruncnormal 100)*(44/12)
   StandDead tC ha
                             = mean(D rtruncnormal 100)
35
   StandDead tCO2 ha
                             = mean(D rtruncnormal 100)*(44/12)
   LyingDead tC ha
                             = mean(E rtruncnormal 100)
37
                             = mean(E_rtruncnormal_100)*(44/12)
   LyingDead_tCO2_ha
   Litter_tC_ha
                             = mean(F_rtruncnormal_100)
39
                             = mean(F_rtruncnormal_100)*(44/12)
   Litter_tCO2_ha
                             = mean(G_rtruncnormal_100)
   Sum_wo_Soil_tC_ha
41
   Sum_wo_Soil_tCO2_ha
                             = mean(G_rtruncnormal_100)*(44/12)
42
   Soil_tC_ha
                             = mean(H_rtruncnormal_100)
43
   Soil_tCO2_ha
                             = mean(H_rtruncnormal_100)*(44/12)
44
45
   CarbonStocks_MC_R_df <- data.frame(</pre>
46
                              = c("tC/ha", "tCO2/ha"),
     Units
47
     `AG Tree`
                              = c(AG tree tC ha, AG tree tCO2 ha),
48
     `BG Tree`
                              = c(BG_tree_tC_ha, BG_tree_tCO2_ha),
49
                              = c(Saplings_tC_ha, Saplings_tCO2_ha),
     `Saplings`
50
                              = c(StandDead_tC_ha, StandDead_tCO2_ha),
     `Standing Dead`
                              = c(LyingDead_tC_ha, LyingDead_tCO2_ha),
     `Lving Dead`
52
                              = c(Litter_tC_ha, Litter_tCO2_ha),
     `Litter`
     `Sum w/o Soil`
                              = c(Sum_wo_Soil_tC_ha, Sum_wo_Soil_tCO2_ha),
54
     `Soil`
                              = c(Soil_tC_ha, Soil_tCO2_ha)
56
   Degradation data
   A_MEAN = DegradationEF$ldf[4]
         = DegradationEF$1df[5]
   B_MEAN = DegradationEF$wood_density[4]
         = DegradationEF$wood_density[5]
   C MEAN = DegradationEF$lif[4]
        = DegradationEF$lif[5]
   D MEAN = DegradationEF$forestry infrastructure[7]
        = DegradationEF$forestry_infrastructure[8]
   E MEAN = DegradationEF$mining[7]
        = DegradationEF$mining[8]
   F MEAN = DegradationEF$mining infrastructure[7]
11
   F_SD = DegradationEF$mining_infrastructure[8]
   G_MEAN = DegradationEF$infrastructure[7]
13
   G_SD = DegradationEF$infrastructure[8]
15
   # 100 simulations sampling 1 observation each iteration.
16
   A rtruncnormal 100 = truncnorm::rtruncnorm(n=100,0,Inf,A MEAN, A SD)
17
   B_rtruncnormal_100 = truncnorm::rtruncnorm(n=100,0,Inf,B_MEAN, B_SD)
18
   C_rtruncnormal_100 = truncnorm::rtruncnorm(n=100,0,Inf,C_MEAN, C_SD)
19
   D_rtruncnormal_100 = truncnorm::rtruncnorm(n=100,0,Inf,D_MEAN, D_SD)
20
   E_rtruncnormal_100 = truncnorm::rtruncnorm(n=100,0,Inf,E_MEAN, E_SD)
21
   F rtruncnormal 100 = truncnorm::rtruncnorm(n=100,0,Inf,F MEAN,F SD)
   G_rtruncnormal_100 = truncnorm::rtruncnorm(n=100,0,Inf,G_MEAN, G_SD)
23
24
```

```
LDF EF tCO2 m2
                              = mean(A rtruncnormal 100)
   WD EF tCO2 m2
                              = mean(B rtruncnormal 100)
26
                              = mean(C rtruncnormal 100)
   LIF EF tCO2 km
27
                              = mean(D rtruncnormal 100)
   ForInfr_EF_tCO2_ha
                              = mean(E rtruncnormal 100)
   Mining_EF_tCO2_ha
29
   MiningInfr EF tCO2 ha
                              = mean(F rtruncnormal 100)
   Infrastructure EF tCO2 ha= mean(G rtruncnormal 100)
31
   df_logging <- data.frame(</pre>
33
      component = c("LDF", "Wood Density of Harvest", "LIF (Skid Trails)"),
34
                = c("per m3", "per m3", "per km"),
35
                = c(LDF_EF_tCO2_m2, WD_EF_tCO2_m2, LIF_EF_tCO2_km)
     tco2
36
37
38
   df_degrading <- data.frame(degrading_activity = c(</pre>
39
        "Forestry infrastructure", "Mining (medium & large scale)",
40
        "Mining infrastructure", "Infrastructure"),
41
     ef tco2 ha = c(ForInfr EF tCO2 ha, Mining EF tCO2 ha,
42
       MiningInfr_EF_tCO2_ha, Infrastructure_EF_tCO2_ha)
43
44
45
   max rows <- max(nrow(df logging), nrow(df degrading))</pre>
46
                         = rep(NA, max_rows - nrow(df_logging))
   logging_nas
                         = rep(NA, max_rows - nrow(df_degrading))
   degrading nas
48
   Degradation MC R df = data.frame(
     Component
                         = c(df logging$component, logging nas),
50
     Unit
                         = c(df_logging$unit, logging_nas),
51
                         = c(df_logging$tco2, logging_nas),
52
     Degrading_Activity= c(df_degrading$degrading_activity, degrading_nas),
53
                         = c(df degrading$ef tco2 ha, degrading nas)
     EF tCO2 ha
54
     )
55
```

Activity data

For purpose of saving space, columns and cells C1-M8 were implemented in the full script but omitted in the final render with chunk settings applied with echo-F. These, and can be located in the associated markdown.Rmd file provided with this PDF.

```
A1 MEAN = ActivityData$x2011[1]
          = ActivityData$x2011[17]
   A1 SD
   A2_MEAN = ActivityData$x2011[2]
          = ActivityData$x2011[18]
   A3 MEAN = ActivityData$x2011[3]
           = ActivityData$x2011[19]
   A3 SD
   A4_MEAN = ActivityData$x2011[4]
           = ActivityData$x2011[20]
   A5_MEAN = ActivityData$x2011[5]
   A5_SD
           = ActivityData$x2011[21]
10
   A6_MEAN = ActivityData$x2011[6]
           = ActivityData$x2011[22]
12
   A7_MEAN = ActivityData$x2011[7]
   A7 SD
           = ActivityData$x2011[23]
14
   A8_MEAN = ActivityData$x2011[8]
          = ActivityData$x2011[24]
   A8 SD
16
```

```
B1 MEAN = ActivityData$x2012[1]
          = ActivityData$x2012[17]
19
   B2 MEAN = ActivityData$x2012[2]
20
          = ActivityData$x2012[18]
   B2 SD
   B3 MEAN = ActivityData$x2012[3]
22
           = ActivityData$x2012[19]
   B4_MEAN = ActivityData$x2012[4]
24
   B4 SD
           = ActivityData$x2012[20]
   B5 MEAN = ActivityData$x2012[5]
26
   B5_SD
           = ActivityData$x2012[21]
   B6_MEAN = ActivityData$x2012[6]
28
   B6_SD
           = ActivityData$x2012[22]
   B7_MEAN = ActivityData$x2012[7]
30
          = ActivityData$x2012[23]
31
   B8_MEAN = ActivityData$x2012[8]
32
   B8 SD
           = ActivityData$x2012[24]
33
   A1 rtruncnormal 100 = truncnorm::rtruncnorm(n=100, 0, Inf, A1 MEAN, A1 SD)
35
   A2_rtruncnormal_100 = truncnorm::rtruncnorm(n=100, 0, Inf, A2_MEAN, A2_SD)
36
   A3 rtruncnormal 100 = truncnorm::rtruncnorm(n=100, 0, Inf, A3 MEAN, A3 SD)
37
   A4_rtruncnormal_100 = truncnorm::rtruncnorm(n=100, 0, Inf, A4_MEAN, A4_SD)
   A5 rtruncnormal 100 = truncnorm::rtruncnorm(n=100, 0, Inf, A5 MEAN, A5 SD)
39
   A6_rtruncnormal_100 = truncnorm::rtruncnorm(n=100, 0, Inf, A6_MEAN, A6_SD)
   A7_rtruncnormal_100 = truncnorm::rtruncnorm(n=100, 0, Inf, A7_MEAN, A7_SD)
41
   A8 rtruncnormal 100 = truncnorm::rtruncnorm(n=100, 0, Inf, A8 MEAN, A8 SD)
43
   B1_rtruncnormal_100 = truncnorm::rtruncnorm(n=100, 0, Inf, B1_MEAN, B1_SD)
   B2 rtruncnormal_100 = truncnorm::rtruncnorm(n=100, 0, Inf, B2_MEAN, B2_SD)
45
   B3 rtruncnormal_100 = truncnorm::rtruncnorm(n=100, 0, Inf, B3_MEAN, B3_SD)
   B4_rtruncnormal_100 = truncnorm::rtruncnorm(n=100, 0, Inf, B4_MEAN, B4_SD)
47
   B5_rtruncnormal_100 = truncnorm::rtruncnorm(n=100, 0, Inf, B5_MEAN, B5_SD)
   B6_rtruncnormal_100 = truncnorm::rtruncnorm(n=100, 0, Inf, B6_MEAN, B6_SD)
40
   B7_rtruncnormal_100 = truncnorm::rtruncnorm(n=100, 0, Inf, B7_MEAN, B7_SD)
50
   B8_rtruncnormal_100 = truncnorm::rtruncnorm(n=100, 0, Inf, B8_MEAN, B8_SD)
52
   # --- Simulation Estimates ---
53
   A1 = mean(A1 rtruncnormal 100)
54
   A2 = mean(A2_rtruncnormal_100)
   A3 = mean(A3 rtruncnormal 100)
56
   A4 = mean(A4_rtruncnormal_100)
   A5 = mean(A5 rtruncnormal 100)
58
   A6 = mean(A6 rtruncnormal 100)
   A7 = mean(A7 rtruncnormal 100)
60
   A8 = mean(A8_rtruncnormal_100)
61
62
   B1 = mean(B1_rtruncnormal_100)
   B2 = mean(B2 rtruncnormal 100)
64
   B3 = mean(B3_rtruncnormal_100)
   B4 = mean(B4_rtruncnormal_100)
   B5 = mean(B5_rtruncnormal_100)
   B6 = mean(B6_rtruncnormal_100)
   B7 = mean(B7_rtruncnormal_100)
   B8 = mean(B8 rtruncnormal 100)
```

Below, we presented organize our results into a table and compare with absolute input values (CarbonStocks), and Monte Carlo estimates generated with SimVoi (CarbonStocks (MC)). For external comparisons, we have also saved these results in a new excel tab called "CarbonStocks (MC-R)".

```
1   CarbonStocks_MC_R = flextable(head(CarbonStocks_MC_R_df[, 1:8])) |>
2   fontsize(size = 8, part = "all")
3   CarbonStocks_MC_R

5   Degradation_MC_R = flextable(head(Degradation_MC_R_df[, 1:8])) |>
6   fontsize(size = 8, part = "all")
7   Degradation_MC_R
```

Table 4: Results of Monte Carlo simulations of Carbon Stocks tabsheet using R

```
CarbonStocks_MC_R = flextable(CarbonStocks_MC_R_df) |>
width(width = 1) |> fit_to_width(max_width = 6) |>
colformat_double(big.mark = ",", digits = 1, na_str = "N/A")
CarbonStocks_MC_R
```

Units	AG.Tree	BG.Tree	Saplings	Standing.Dead	Lying.Dead	Litter	Sum.w.o.Soil	Soil
tC/ha	210.9	48.5	3.9	4.5	11.0	3.4	267.1	82.7
tCO2/ha	773.4	177.9	14.3	16.5	40.3	12.6	979.5	303.1

Table 5: Results of Monte Carlo simulations of CarbonStocks tabsheet using SimVoi

Units	AG Tree	BG Tree	Saplings	Standing Dead	Lying Dead	Litter	Sum w/o Soil	Soil
tC/ha	181.1	65.0	3.5	7.3	17.1	3.7	277.7	60.6
tCO2/ha	664.2	238.2	12.8	26.9	62.6	13.7	1,018.4	222.3

Table 6: Results of Monte Carlo simulations of DegradationEFs tabsheet using R

Component	Unit	tCO2	Degrading_Activity	EF_tCO2_ha
LDF	per m3	4.4	Forestry infrastructure	10.6
Wood Density of Harvest	per m3	1.5	Mining (medium & large scale)	9.4
LIF (Skid Trails)	per km	167.6	Mining infrastructure	11.1
		N/A	Infrastructure	11.4

Table 7: Results of Monte Carlo simulations of Degradation EFs tabsheet using SimVoi

Component	Unit	t CO2	Degrading Activity	EF (tCO2/ha)
LDF	per m ³	4.54	Forestry infrastructure	7.2
Wood Density of timber harvested	per m³	1.39	Mining (med & large scale)	7.2
LIF (Skid Trails)	per km	185.88	Mining infrastructure	7.2
			Infrastructure	7.2

Table 8: Results of Monte Carlo simulations of Activity Data tabsheet using R

Drivers	units	X2011	X2012	X2013	X2014	X2015
Forestry infrastructure	ha	185	242	328	205	322
Agriculture	ha	41	429	422	810	381
Mining (med & large scale)	ha	7,374	13,667	11,321	10,455	6,822
Mining infrastructure	ha	N/A	N/A	N/A	N/A	N/A
Infrastructure	ha	119	127	344	140	218
Settlements	ha	N/A	N/A	24	70	8
Fire-Biomass burning	ha	46	187	95	259	1,541
Shifting Cultivation	ha	N/A	N/A	N/A	N/A	N/A

Table 9: Results of Monte Carlo simulations of Activity Data tabsheet using SimVoi

Drivers	units	2011	2012	2013	2014	2015	2016	2017	2018	2019	2020	2021	2022
Forestry infrastructure		225	194	270	229	325	288	224	244	228	185	215	83
Agriculture		36	384	463	882	436	401	487	291	261	126	436	305
Mining (med & large scale)		8,835	13,157	7,687	12,583	7,673	7,082	6,619	6,295	5,518	6,980	7,303	6,163
Infrastructure	ha	117	121	331	143	182	236	196	72	33	188	101	117
Settlements		-	-	25	73	7	7	7	9	29	53	110	210
Fire-Biomass burning		53	188	89	301	1,833	1,553	460	681	6,194	3,364	221	365
Shifting Cultivation								451	361	509	478	456	109
Deforestation		9,266	14,044	8,863	14,210	10,457	9,568	7,993	7,592	12,772	11,373	8,842	7,353
Logging - harvest volume	m3	608,730	585,108	624,287	759,684	655,406	500,788	533,106	546,242	521,172	545,355	547,516	547,517
Logging - skid trail length	km	2,302	2,212	2,360	2,872	2,478	1,893	2,016	2,065	1,971	2,062	2,070	2,070
Illegal logging	m3	2776	2,306	2,371	2,836	1,505	2,249	2,706	3,719	2,149	1,281	1,281	1,281
Mining and Infrastructure	ha						36,647	31,919	28,185	23,028	22,795	26,651	26,651

Replication results

In the following chunk, we compute the final uncertainty estimates for the simulated emission reductions (GHG ER) using a truncated normal distribution. With 10,000 simulation trials, we calculate key statistics, mean, standard deviation, and mean standard error, along with distributional percentiles. By extracting the 5th and 95th percentiles, we form a 90% confidence interval (CI) from which we derive the margin of error (ME) and its corresponding percentage error relative to the mean. These computed metrics replicate the SimVoi univariate summary and provide a statistical summary of the uncertainty associated with the emissions reduction estimates.

```
# Emission Reductions (MC-R)
ER_values <- c(7715885, 10371977, 10040723, 6358705, 7174999, 6977178, 9223423, 7299024)
ER_mean_emp <- mean(ER_values)</pre>
ER_sd_emp
            <- sd(ER_values)
n sim <- 10000
\# Simulate a truncated normal distribution \mho compute stats
sim_ER <- rtruncnorm(n = n_sim, a = 0, b = Inf, mean = ER_mean_emp, sd = ER_sd_emp)
sim_mean <- mean(sim_ER)</pre>
sim sd
       <- sd(sim_ER)
         <- sim_sd / sqrt(length(sim_ER))
sim_skew <- moments::skewness(sim_ER)</pre>
sim_quant <- quantile(sim_ER, probs = c(0, 0.25, 0.5, 0.75, 1))
lower90 <- quantile(sim_ER, probs = 0.05)</pre>
upper90 <- quantile(sim_ER, probs = 0.95)</pre>
ci90 <- upper90 - lower90
```

```
ME < - ci90 / 2
pct_error <- ME / sim_mean * 100</pre>
ER_summary <- data.frame(</pre>
  Metric = c("Mean", "St. Dev.", "Mean St. Error", "Skewness",
             "Minimum", "1st Quartile", "Median", "3rd Quartile", "Maximum",
             "5th Percentile", "95th Percentile", "90% CI", "Margin of Error", "% Error"),
  Value = c(round(sim mean),
                                   round(sim_sd),
             round(sim se),
                                   round(sim skew, 3),
             round(sim_quant[1]), round(sim_quant[2]),
             round(sim_quant[3]), round(sim_quant[4]),
             round(sim_quant[5]), round(lower90),
             round(upper90), round(ci90),
             round(ME), round(pct_error, 2))
  )
```

Table 10: Simulated Univariate Summary of Emission Reductions (GHG ER)

Metric	Value
Mean	8,149,938.000
St. Dev.	1,511,217.000
Mean St. Error	15,112.000
Skewness	0.023
Minimum	2,251,188.000
1st Quartile	7,128,275.000
Median	8,135,566.000
3rd Quartile	9,163,819.000
Maximum	13,632,541.000
5th Percentile	5,689,744.000
95th Percentile	10,628,701.000
90% CI	4,938,957.000
Margin of Error	2,469,479.000
% Error	30.300

Distribution analysis

Distribution analysis is a critical to ensuring Monte Carlo simulations accurately reflect the empirical characteristics of input data. Visual and statistical checks, such as normality tests (Shapiro, Sandford & Wilk, 1965) and distribution plots (Figures 9–16), verify data shape, spread, skewness, and outliers. Overlooking this step risks masking or exaggerating inherent biases, thereby undermining reliability of estimates. Figures 1–8 display the simulated distributions, while Figures 9–16 show the original data distributions. Discrepancies between them highlight opportunities for reducing uncertainty. We recommend to refine the current simulation approach by incorporating data-specific models, such as simulations fitted with log-normal and beta distributions.

```
# Distribution of Monte Carlo derived estimates
hist(A_rtruncnormal_100, freq=F, main="Aboveground Tree Biomass (tC/ha)")
```

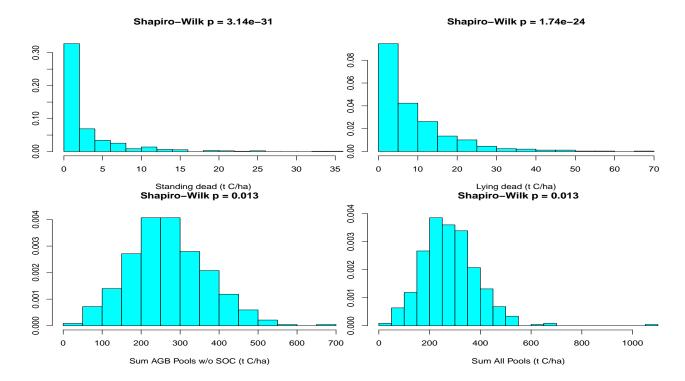
```
hist(B_rtruncnormal_100, freq=F, main="Belowground Tree Biomass (tC/ha)")
hist(C_rtruncnormal_100, freq=F, main="Saplings Biomass (tC/ha)")
hist(D_rtruncnormal_100, freq=F, main="Standing Dead Wood (tC/ha)")
hist(E_rtruncnormal_100, freq=F, main="Lying Dead Wood (tC/ha)")
hist(F_rtruncnormal_100, freq=F, main="Litter Biomass (tC/ha)")
hist(G_rtruncnormal_100, freq=F, main="Sum Pools w/o Soil (tC/ha)")
hist(H_rtruncnormal_100, freq=F, main="Soil Biomass (tC/ha)")
                     Aboveground Tree Biomass (tC/ha)
                                                                                         Belowground Tree Biomass (tC/ha)
     0.20
     0.15
Density
                                                                    Density
    0.10
     0.02
     0.00
                                                                         0
           0
                      2
                                            6
                                                       8
                                                                  10
                                                                              1.2
                                                                                        1.3
                                                                                                  1.4
                                                                                                             1.5
                                                                                                                       1.6
                                                                                                                                 1.7
                         A_rtruncnormal_100
Saplings Biomass (tC/ha)
                                                                                            B_rtruncnormal_100
Standing Dead Wood (tC/ha)
    0.015
                                                                         90.0
     0.010
Density
                                                                         0.04
                                                                    Density
    0.005
                                                                         0.02
    0.000
                                                                         0.00
                                                                                       5
          100
                 120
                                 160
                                         180
                                                 200
                                                         220
                                                                240
                                                                                                      15
                                                                                                              20
                                                                                                                      25
                                                                                                                              30
                                                                                                                                      35
                         C_rtruncnormal_100
Lying Dead Wood (tC/ha)
                                                                                               D_rtruncnormal_100
Litter Biomass (tC/ha)
     90.0
                                                                         0.0
     0.04
                                                                    Density
Density
                                                                         0.02
    0.02
    0.00
                                                                         0.00
                                                                                        5
          0
                  5
                          10
                                  15
                                          20
                                                  25
                                                          30
                                                                  35
                                                                               0
                                                                                                 10
                                                                                                          15
                                                                                                                   20
                                                                                                                            25
                                                                                                                                      30
                         E_rtruncnormal_100
Sum Pools w/o Soil (tC/ha)
                                                                                                F_rtruncnormal_100
Soil Biomass (tC/ha)
    90.0
                                                                         0.008
     0.04
Density
                                                                    Density
                                                                         0.004
     0.02
                                                                         0.000
    0.00
                  5
                          10
                                  15
                                          20
                                                          30
                                                                                           50
                                                                                                       100
                                                                                                                    150
                                                                                                                                 200
                              G_rtruncnormal_100
                                                                                                   H_rtruncnormal_100
```

4

5

Figures 1-8: Distribution analysis of simulated estimates of carbon stock variables

```
inventory dataset = "./data/art/AllBiomassDataCombined Master September2019 ANALYSIS 2021 DescStat His
   tree_data = readxl::read_excel(inventory_dataset, "Distribution R") |>
     janitor::clean_names() |> mutate(across(where(is.numeric), ~ round(.x, 1)))
3
   sw trees agb = stats::shapiro.test(tree data$trees agb t c ha)
5
   sw_trees_bgb = stats::shapiro.test(tree_data$trees_bgb_t_c_ha)
   sw saplings = stats::shapiro.test(tree data$saplings t c ha)
   sw_litter = stats::shapiro.test(tree_data$litter_t_c_ha)
   sw_standing_dead = stats::shapiro.test(tree_data$standing_dead_t_c_ha)
   sw lying dead = stats::shapiro.test(tree data$lying dead t c ha)
10
   sw agb pools = stats::shapiro.test(tree data$sum agb pools no soil t c ha)
11
   sw all pools = stats::shapiro.test(tree data$sum all pools t c ha)
13
   MASS::truehist(tree_data$trees_agb_t_c_ha, xlab = "Above-ground biomass (t C/ha)",
14
     main = sprintf("Shapiro-Wilk p = %.3g", sw_trees_agb$p.value))
15
   MASS::truehist(tree data$trees bgb t c ha, xlab = "Below-ground biomass (t C/ha)",
16
     main = sprintf("Shapiro-Wilk p = %.3g", sw_trees_bgb$p.value))
   MASS::truehist(tree_data$saplings_t_c_ha, xlab = "Sapling (t C/ha)",
18
     main = sprintf("Shapiro-Wilk p = %.3g", sw_saplings$p.value))
19
   MASS::truehist(tree_data$litter_t_c_ha, xlab = "Litter (t C/ha)",
20
     main = sprintf("Shapiro-Wilk p = %.3g", sw_litter$p.value))
   MASS::truehist(tree_data$standing_dead_t_c_ha, xlab = "Standing dead (t C/ha)",
22
     main = sprintf("Shapiro-Wilk p = %.3g", sw_standing_dead$p.value))
23
   MASS::truehist(tree_data$lying_dead_t_c_ha, xlab = "Lying dead (t C/ha)",
24
     main = sprintf("Shapiro-Wilk p = %.3g", sw_lying_dead$p.value))
   MASS::truehist(tree_data$sum_agb_pools_no_soil_t_c_ha, xlab = "Sum AGB Pools w/o SOC (t C/ha)",
26
     main = sprintf("Shapiro-Wilk p = %.3g", sw_agb_pools$p.value))
27
   MASS::truehist(tree_data$sum_all_pools_t_c_ha, xlab = "Sum All Pools (t C/ha)",
28
     main = sprintf("Shapiro-Wilk p = %.3g", sw_agb_pools$p.value))
                    Shapiro-Wilk p = 0.0031
                                                                     Shapiro-Wilk p = 0.0071
                                                    0.020
   0.00
                                                    0.010
   0.002
                                                    0.00
   0.000
                                              500
                                                               20
                                                                                                120
                    Above-ground biomass (t C/ha)
                                                                    Below-ground biomass (t C/ha)
                  Shapiro-Wilk p = 1.06e-22
                                                                    Shapiro-Wilk p = 3.01e-35
   0.25
                                                    5
   0.20
   0.15
                                                    9
   0.10
                                                    0.5
   0.05
                                                    0.0
        O
                      10
                             15
                                    20
                                           25
                                                  30
                                                         o
                                                                                               8
                                                                          Litter (t C/ha)
                        Sapling (t C/ha)
```



Figures 9-16: Distribution analysis of carbon stock input variables

In addition, these distributional visualizations may above offer auditors useful diagnostic tools, enabling rapid identification and characterization of biases commonly encountered in biomass data. Such diagrams help auditors efficiently assess the statistical approaches implemented to monitor and manage uncertainty in the project's data.

To guide practitioners in appropriate simulation designs, the following two tables present findings from a rapid literature review of Monte Carlo methods in forestry and REDD+ contexts (Annex II).

Table 9: Continuous data distributions, example cases $\mathfrak E$ equations used in Monte Carlo simulations.

Distribution	Statistical Use Cases	PDF
Normal	Symmetric, bell-shaped distribution used for modeling continuous variables: biomass/ha	$f(x) = \frac{1}{\sigma\sqrt{2\pi}} \exp\left(-\frac{(x-\mu)^2}{2\sigma^2}\right)$
Lognormal	Right-skewed distribution suitable for variables constrained to positive values (e.g., emission rates).	$f(x) = \frac{1}{x\sigma\sqrt{2\pi}}\exp\left(-\frac{(\ln x - \mu)^2}{2\sigma^2}\right)$
Exponential	Models waiting times between independent events, such as forest fire occurrences or logging events.	$f(x) = \lambda e^{-\lambda x}, \\ x \ge 0$
Cont. Uniform	Assumes all values in an interval [a, b] are equally likely; useful for random spatial sampling in forests.	$f(x) = \frac{1}{b-a},$ $a < x < b$
Chi-Square	Often used in goodness-of-fit tests to evaluate model accuracy in biomass estimation.	$f(x) = \frac{1}{2^{k/2} \Gamma(k/2)} x^{\frac{k}{2} - 1} e^{-x/2}, x > 0$
t-Distribution	Suitable for small sample sizes with unknown population stdev (e.g., limited forest carbon data).	$f(x) = \frac{\Gamma\left(\frac{v+1}{2}\right)}{\sqrt{\nu\pi}\Gamma\left(\frac{v}{2}\right)} \left(1 + \frac{x^2}{v}\right)^{\frac{v+1}{2}}$
Gamma	Models positively skewed data, such as biomass growth rates or carbon accumulation over time.	$f(x) = \frac{x^{k-1}e^{-x/\theta}}{\theta^k \Gamma(k)}$
Weibull	Flexible distribution used in reliability analysis, e.g., modeling tree mortality.	$f(x) = \frac{k}{\lambda} \left(\frac{x}{\lambda}\right)^{k-1} e^{-(x/\lambda)^k}$

Table 10: Discrete data distributions, example cases $\mathfrak E$ equations used in Monte Carlo simulations.

Distribution	Statistical Use Cases	PMF
Bernoulli	Binary outcome probability, e.g., presence/absence of deforestation in an area.	$P(X = x) = p^{x}(1-p)^{1-x},$ $x \in \{0,1\}$
Binomial	Probability of fixed #no. of successes over n Bernoulli trials, e.g., no. of heads in ten coin flips.	$P(X = k) = \binom{n}{k} p^k (1 - p)^{n-k},$ k = 0, 1,, n
Poisson	Models counts of independent events within an interval, e.g., number of wildfire incidents per year.	$P(X = k) = \frac{\lambda^k e^{-\lambda}}{k!},$ k = 0,1,2,
Geometric	Models #no. of trials until the first success, e.g., number of inspections until detecting deforestation.	$P(X = k) = (1 - p)^{k-1}p,$ k = 1, 2,
Negative Binomial	Counts #no. failures until r successes occur, treats overdispersed or repeated deforestation detections.	$P(X = k) = {k + r - 1 \choose k} (1 - p)^r p^k, k = 0,1,2,$
Discrete Uniform	Assumes outcome in a finite set is equally likely, e.g., random sampling of inventory across a forest.	$P(X = x) = \frac{1}{n},$ x = 1,2,,n

Discrete distributions model countable events in forestry, such as deforestation counts, wildfire occurrences, or logged trees, using tools like the Binomial, Poisson, or Negative Binomial distributions to improve prediction accuracy and uncertainty assessments. For example, a Poisson distribution can enhance precision in estimating deforestation emissions from illegal logging.

Continuous distributions apply to variables that assume any value within a range, such as tree heights, carbon stock densities, or biomass. Common models, including Normal, Lognormal, Weibull, and Gamma distributions, capture ecological variability effectively; for instance, a Lognormal distribution often provides more reliable biomass estimates for right-skewed data.

The core mathematical concepts are the Probability Mass Function (PMF) for discrete data and the Proba-

bility Density Function (PDF) for continuous data. Accurate use of PMFs and PDFs is essential in Monte Carlo simulations, as they underpin random sampling processes that directly influence the reliability of uncertainty estimates. Rigorous selection of these functions enhances biomass and emissions estimates, reduces uncertainty, and supports the credibility of REDD+ reporting (Morgan & Henrion, 1990; IPCC, 2019; ART, 2021).

Early exploratory data analysis, including statistical normality tests and visual assessments (histograms, kernel density plots, Q-Q plots), is recommended to diagnose data distributions, optimize model selection, and reduce audit findings, ultimately improving the financial and environmental outcomes of national REDD+monitoring programs.

Recommendations

- Regarding 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 formula applied in Guyana's verified workbook uses a minimum value of 0. Considering the levels of quantitative variance between input values in the "CarbonStocks" tabsheet and SimVoisimulated estimates in "CarbonStocks (MC)" tabsheet, we may assume that approximately less than ~100 simulations were completed before values were reached "between MinValue and MaxValue". It is useful to consider that with increasing numbers of simulations the closer the mean values generated should be to absolute input values. Therefore, in order to test this assumption we simply ran 10,000 simulations, which generated mean estimates of within 2 tC ha-1 of input values (205 tC ha-1) 10 times of 10 reruns. By reducing the number of iterations per simulation, we confirm our assumptions that the current SimVoi setup provided estimates from ~<100 simulations.
- We recommend increasing the number of simulation iterations from 100 to a higher number, in order to reduce variance in and between simulated estimates, which thereby provide more robust uncertainty quantification.
- It is advised to consider performing sensitivity analyses in order to identify which input parameters
 contribute most to overall uncertainty, thereby informing targeted data collection and model refinement
 efforts.
- Furthermore, distribution analysis confirms that our simulation outputs approximate the expected statistical distributions of the underlying data. In the next phase, we recommend incorporating the observed distributional characteristics into the Monte Carlo simulation framework. This adjustment, using distribution-specific models (e.g., log-normal, beta, etc.), is essential for reducing uncertainty and ensuring that our estimates carry statistical significance.

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Annex I: SimVoi replication in base-R

Using the following base functions from R avoids need for any package installations and limits risk of bugs or common problems with R versioning arising in the workflow. However, the base functions below are a little clunkier and may require additional guidance. Regarding R and package versions, please consult the Annex II which presents record of the current runtime used in this analysis.

```
randtruncnormal_sim_10000 <- rnorm(n=10,mean=MEAN,sd=SD)
hist(randtruncnormal_sim_10000, freq=F)
AG_Tree_tC_ha = mean(randtruncnormal_sim_10000)
AG_Tree_tC02_ha = AG_Tree_tC_ha*(44/12)
AG_Tree_tC_ha
AG_Tree_tC02_ha
#curve(dnorm(x, mean=MEAN, sd=SD), from=0, to=450, add=T, col="red")</pre>
```

```
# 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)
```

Annex II: Review of Monte Carlo methods in REDD+

Table A.2: Search parameters, resource scope, and objectives informing search

REDD+ ³ ADD	MC Application Uncertainty of SAAB estimate	Region Rondônia, Brazil	Key Findings Estimated ± 20% measurement error in SAAB using Monte Carlo simulations; emphasized large trees' role in biomass.
ADD	AGB Uncertainty	Kenya, Mozambique	Assessed mixed-effects models in estimating mangrove biomass.
ADD	Blanket uncertainty propagation	Ghana	AGB prediction error >20%; addressed error propagation from trees to pixels in remote sensing.
ADD	Plot-based uncertainty	New Zealand	Cross-plot variance greatest magnitude of uncertainty

³1. ADD: Avoided deforestation degradation, IFM: Improved forest management, JNR: Jurisdictional nested REDD+

JNR	Multi-scale AGB uncertainty modeling	Minnesota, USA	Cross-scale tests showing effects of spatial resolution on AGB
N/A	Allometric uncertainty modeling	Panama	uncertainty. Allometric models identified as largest source of biomass estimation error.
ADD	Sampling and allometric uncertainty	Tapajos Nat Forest, Brazil	Significance of allometric models on uncertainty of root biomass, 95% CI, 21 plots.
ADD	Uncertainty of volume estimates	Santa Catarina, Brazil	Negligible effects of residual uncertainty on large-area estimates
N/A	Uncertainty metrics in model selection	Oregon, USA	Uncertainty estimates call for local validation or new local model development
ADD	AGB model uncertainty	French Guiana	AGB sub-model errors dominate uncertainty; height and wood-specific gravity errors are minor but can cause bias.
IFM	Emission factor uncertainty	Central Africa	Model selection is the largest error source (40%); weighting models reduces uncertainty in emission factors.
NA	Uncertainty in ecosystem nutrient estimate	New Hampshire, USA	Identified 8% uncertainty in nitrogen budgets, mainly from plot variability (6%) and allometric errors (5%).

Annex III: Runtime snapshot

devtools::session_info()

```
- Session info -----
setting value
version R version 4.3.0 (2023-04-21)
      macOS 15.4.1
system aarch64, darwin20
      X11
ui
language (EN)
collate en_US.UTF-8
ctype en_US.UTF-8
tz
      America/Vancouver
      2025-05-05
date
pandoc 3.6.1 @ /usr/local/bin/ (via rmarkdown)
- Packages ------
```

package	*	version	date (UTC)	lih	sourc	۵.	
animation		2.7	2021-10-07	[1]			4.3.3)
askpass		1.2.1	2024-10-04	[1]	CRAN		4.3.3)
assertthat		0.2.1	2019-03-21	[1]	CRAN		4.3.0)
backports		1.5.0	2024-05-23	[1]	CRAN		4.3.3)
BIOMASS	.	2.2.3	2024 03 23	[1]	CRAN		4.3.3)
boot	т	1.3-31	2024-08-28	[1]	CRAN		4.3.3)
	*	1.0.7	2024-00-26	[1]	CRAN		4.3.3)
broom							•
c2z	*	0.2.0 1.1.0	2023-08-10	[1] [1]	CRAN CRAN		4.3.0)
cachem	.1.		2024-05-16				4.3.3)
caret	*	7.0-1	2024-12-10	[1]	CRAN		4.3.3)
cellranger		1.1.0	2016-07-27	[1]	CRAN		4.3.0)
chromote		0.4.0	2025-01-25	[1]	CRAN		4.3.3)
class		7.3-23	2025-01-01	[1]	CRAN		4.3.3)
classInt		0.4-11	2025-01-08	[1]	CRAN		4.3.3)
cli		3.6.3	2024-06-21	[1]	CRAN		4.3.3)
codetools		0.2-20	2024-03-31	[1]	CRAN		4.3.1)
colorspace		2.1-1	2024-07-26	[1]	CRAN		4.3.3)
data.table		1.16.4	2024-12-06	[1]	CRAN		4.3.3)
dataMaid	*	1.4.1	2021-10-08	[1]	CRAN		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		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)
distill	*	1.6	2023-10-06	[1]	CRAN	(R	4.3.1)
downlit		0.4.4	2024-06-10	[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		4.3.3)
Exact		3.3	2024-07-21	[1]	CRAN	(R	4.3.3)
expm		1.0-0	2024-08-19	[1]	CRAN		4.3.3)
extrafont	*	0.19	2023-01-18	[1]	CRAN	(R	4.3.3)
extrafontdb		1.0					4.3.3)
fastmap		1.2.0	2024-05-15	[1]	CRAN		4.3.3)
flextable	*	0.9.7	2024-10-27	[1]	CRAN		4.3.3)
fontBitstreamVera		0.1.1	2017-02-01	[1]	CRAN		4.3.3)
fontLiberation		0.1.0	2016-10-15	[1]	CRAN		4.3.3)
fontquiver		0.2.1	2017-02-01	[1]	CRAN		4.3.3)
forcats	*	1.0.0	2023-01-29		CRAN		4.3.0)
foreach		1.5.2	2022-02-02	[1]	CRAN		4.3.0)
formatR	*	1.14	2023-01-17	[1]	CRAN		4.3.3)
fs	т	1.6.5	2023 01 17	[1]	CRAN		4.3.3)
furrr		0.3.1	2024 10 30	[1]	CRAN		4.3.0)
future		1.34.0	2024-07-29	[1]	CRAN		4.3.3)
future.apply		1.11.3	2024-10-27	[1]	CRAN		4.3.3)
gdtools		0.4.1	2024-11-04	[1]	CRAN		4.3.3)
generics		0.1.3	2022-07-05	[1]	CRAN		4.3.0)
ggplot2	*	3.5.1	2024-04-23	[1]	CRAN		4.3.1)
gld		2.6.7	2025-01-17	[1]	CRAN	(K	4.3.3)

globals		0.16.3	2024-03-08			
glue		1.8.0	2024-09-30	[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)
latex2exp	*	0.9.6	2022-11-28	[1]	CRAN	(R 4.3.0)
latexpdf	*	0.1.8	2023-12-19	[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)
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		CRAN	
minpack.lm		1.2-4	2023-09-11		CRAN	
modeldata	*	1.4.0	2024-06-19			
ModelMetrics		1.2.2.2	2020-03-17		CRAN	
moments	*	0.14.1			CRAN	
munsell		0.5.1	2024-04-01	[1]		
mvtnorm		1.3-3			CRAN	
nlme		3.1-166	2024-08-14			
nnet		7.3-20	2025-01-01	[1]		
officer	*	0.6.7	2024-10-09			
openssl	-	2.3.1	2025-01-09	[1]		
pander		0.6.6	2025-03-01	[1]		
parallelly		1.41.0	2024-12-18	[1]		
paranie	*	1.2.1	2024 12 18	[1]		
pillar	-,-	1.10.1	2024-03-22	[1]		
prirar pkgbuild		1.4.6	2025-01-07	[1]		
pkgconfig		2.0.3	2019-09-22		CRAN	
Lv2com 12		2.0.0	2010 00 22	ΓŢĴ	OILHIN	(11 4.0.0)

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2024-06-28 [1] CRAN (R 4.3.3)
pkgload
                     1.4.0
                     1.8.9
                                2023-10-02 [1] CRAN (R 4.3.1)
plyr
pROC
                     1.18.5
                                2023-11-01 [1] CRAN (R 4.3.1)
                                2025-01-08 [1] CRAN (R 4.3.3)
                     3.8.5
processx
prodlim
                     2024.06.25 2024-06-24 [1] CRAN (R 4.3.3)
profvis
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                                2024-09-20 [1] CRAN (R 4.3.3)
                     1.3.2
                                2024-11-28 [1] CRAN (R 4.3.3)
promises
                     0.4 - 27
                                2022-06-09 [1] CRAN (R 4.3.0)
proxy
ps
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                   * 1.0.2
purrr
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R6
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rappdirs
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                                2025-01-12 [1] CRAN (R 4.3.3)
Rcpp
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readxl
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                   * 1.1.0
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remotes
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robustbase
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rootSolve
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rstudioapi
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Rttf2pt1
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scales
sessioninfo
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sf
                     1.0 - 21
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shiny
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stringr
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svglite
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                                2024-04-18 [1] CRAN (R 4.3.1)
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                   * 1.2.1
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units
urlchecker
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useful
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3.1.0
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vctrs
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viridisLite
                 0.4.2
                             2023-05-02 [1] CRAN (R 4.3.0)
                             2023-06-26 [1] CRAN (R 4.3.0)
webshot
                * 0.5.5
webshot2
                * 0.1.1
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                             2024-07-22 [1] CRAN (R 4.3.3)
websocket
                 1.4.2
                 3.0.2
                           2024-10-28 [1] CRAN (R 4.3.3)
withr
                           2024-02-19 [1] CRAN (R 4.3.1)
2024-03-21 [1] CRAN (R 4.3.1)
                * 1.1.4
workflows
workflowsets
               * 1.1.0
xfun
                  0.50
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                             2023-12-04 [1] CRAN (R 4.3.1)
xm12
                  1.3.6
                 1.8-4
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xtable
                             2024-07-26 [1] CRAN (R 4.3.3)
yaml
                 2.3.10
yardstick
                * 1.3.1
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                             2024-01-27 [1] CRAN (R 4.3.1)
zip
                   2.3.1
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- [1] /Library/Frameworks/R.framework/Versions/4.1-arm64/Resources/library
- [2] /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library

[#]Sys.getenv()

^{2 #.}libPaths()