# Package 'SeedMaker'

July 21, 2025

Title Generate a Collection of Seeds from a Single Seed

Version 1.0.0	
<b>Description</b> A mechanism for easily generating and organizing a collection of seeds from a single seed, which may be subsequently used to ensure reproducibility in processes/pipelines that utilize multiple random components (e.g., trial simulation).	
License MIT + file LICENSE	
Encoding UTF-8	
RoxygenNote 7.3.1	
<b>Suggests</b> devtools (>= 2.4.5), knitr (>= 1.45), rmarkdown (>= 2.25), testthat (>= 3.2.0), usethis (>= 2.2.2)	
<b>Imports</b> checkmate (>= 2.3.0), dplyr (>= 1.1.4), rlang (>= 1.1.2), tibble (>= 3.2.1), tidyr (>= 1.3.0)	
<pre>URL https://github.com/hdhshowalter/SeedMaker</pre>	
<pre>BugReports https://github.com/hdhshowalter/SeedMaker/issues</pre>	
Config/testthat/edition 3	
VignetteBuilder knitr	
NeedsCompilation no	
Author Hollins Showalter [aut, cre], Eli Lilly and Company [cph, fnd]	
Maintainer Hollins Showalter < hollins.showalter@gmail.com>	
Repository CRAN	
<b>Date/Publication</b> 2025-01-17 09:00:02 UTC	
Contents	
<b>−c</b>	2
Index	8

2 seed\_gen

seed\_gen

Generate a vector of seeds

## Description

```
seed_gen() generates a vector of seeds from a single seed.
```

## Usage

```
seed_gen(seed, n, sample_vector = 1:1e+06)
```

## Arguments

```
seed A number, interpreted as an integer, or NULL (see set.seed()).

n The number of seeds to generate.

sample_vector The values from which to sample (without replacement).
```

#### **Details**

```
length(sample\_vector) must be >= n.
```

The values of sample\_vector must be unique, positive integers.

#### Value

An integer vector containing n randomly drawn values from sample\_vector.

## **Examples**

```
seeds1 <- seed_gen(
   seed = 1234,
   n = 10
)

seeds2 <- seed_gen(
   seed = 1234,
   n = 10,
   sample_vector = 1:100
)</pre>
```

seed_maker	Construct a list containing a collection of seeds	

#### Description

seed\_maker() constructs a hierarchical list containing a collection of seeds from a single seed.

### Usage

```
seed_maker(
   seed,
   level_names,
   n_per_level,
   sample_vector = rep(list(rlang::expr(1:1e+06)), length(level_names)),
   index_level = rep(TRUE, length(level_names)),
   flatten = FALSE
)
```

## **Arguments**

seed	A number, interpreted as an integer, or NULL (see set.seed()).
level_names	A character vector, or list of character vectors, indicating the element name(s) to use for each level of the returned list.
n_per_level	An integer vector indicating the number of elements to produce at each level of the returned list.
sample_vector	A list of expressions that are evaluable as integer vectors indicating the values from which to sample (without replacement) at each level of the returned list.
index_level	A logical vector indicating whether to index the root name of a respective level.
flatten	"Flatten" the lowest level of the returned list into a named vector?

#### **Details**

length(level\_names) must be equal to length(n\_per\_level) must be equal equal to length(sample\_vector) must be equal to length(index\_level). When these arguments are greater than length 1, the intermediate seeds used to produce the seeds at the lowest level of the returned list are not included therein. In other words, only the collection of seeds at the lowest list level are returned.

If level\_names is a character vector, each of its elements is used as the root name for the list elements produced at the respective level. See seeds\_list1 and seeds\_list2 in **Examples**.

If level\_names is a list of character vectors:

- The element contained in a length 1 vector is used as the root name for the list elements produced at the respective level.
- Elements contained in a vector that is greater than length 1 are used verbatim as the names for the list elements produced at the respective level. The corresponding value of n\_per\_level must match the length of the vector.

See seeds\_list3, seeds\_list4, and seeds\_list5 in **Examples**.

For each element, i, of sample\_vector:

- length(eval(sample\_vector[[i]])) must be >= n\_per\_level[i].
- The values of eval(sample\_vector[[i]]) must be unique, positive integers.

See seeds\_list6, seeds\_list7, and seeds\_list8 in **Examples**.

Element name indexing applies to root name-provided levels only. Thus, for each element, i, of index\_level, if length(level\_names[[i]]) is greater than 1, the value of index\_level[i] is ignored. Otherwise:

- When index\_level[i] is TRUE, the element names at the respective level are indexed from 1 - n\_per\_level[i].
- When index\_level[i] is FALSE, the element name at the respective level is used verbatim, so long as n\_per\_level[i] = 1, since suppressing element name indexing makes sense only for single-element levels.

See seeds\_list9, seeds\_list10, seeds\_list11, seeds\_list12, and seeds\_list13 in **Examples**.

When flatten is set to TRUE, each set of seeds at the lowest level of the returned list is converted into a named vector (versus remaining in the default structure, which is a list of single value elements). When only a single level is specified (via level\_names, n\_per\_level, sample\_vector, and index\_level all being length 1), a named integer vector is returned instead of a list. See seeds\_flat1, seeds\_flat2, seeds\_flat5, seeds\_flat9, seeds\_flat10, and seeds\_flat11 in **Examples**.

#### Value

When flatten is set to FALSE, a hierarchical list with length(level\_names) levels containing a collection of seeds.

When flatten is set to TRUE, a hierarchical list with length(level\_names) - 1 level(s) containing a collection of seeds (where "0 levels" corresponds to a named integer vector - see **Details**).

## **Examples**

```
seeds_list1 <- seed_maker(
    seed = 1234,
    level_names = c("sim"),
    n_per_level = c(10)
)

seeds_list2 <- seed_maker(
    seed = 1234,
    level_names = c("sim", "dataset"),
    n_per_level = c(10, 15)
)

seeds_list3 <- seed_maker(
    seed = 1234,
    level_names = list("sim", c("component_a", "component_b", "component_c")),</pre>
```

```
n_{per_level} = c(10, 3)
try(seeds_list4 <- seed_maker(</pre>
  seed = 1234,
  level_names = list("sim", c("component_a", "component_b", "component_c")),
  n_{per_level} = c(10, 4)
))
seeds_list5 <- seed_maker(</pre>
  seed = 1234,
  level_names = list(
    "sim",
    c("component_a", "component_b", "component_c"),
    "dataset"
  ),
 n_per_level = c(10, 3, 5)
)
seeds_list6 <- seed_maker(</pre>
  seed = 1234,
  level_names = c("sim"),
 n_{per_level} = c(10),
  sample\_vector = list(1:100)
)
seeds_list7 <- seed_maker(</pre>
  seed = 1234,
  level_names = c("sim", "dataset"),
 n_{per_{level}} = c(10, 15),
  sample_vector = list(1:100, 1:1000)
)
seeds_list8 <- seed_maker(</pre>
  seed = 1234,
  level_names = list(
    "sim",
    c("component_a", "component_b", "component_c"),
    "dataset"
  ),
  n_{per_level} = c(10, 3, 5),
  sample_vector = list(1:100, 1:1000, 1:10000)
)
seeds_list9 <- seed_maker(</pre>
  seed = 1234,
  level_names = c("sim"),
  n_per_level = c(1)
)
try(seeds_list10 <- seed_maker(</pre>
  seed = 1234,
  level_names = c("sim"),
```

```
n_{per_level} = c(10),
  index_level = FALSE
))
seeds_list11 <- seed_maker(</pre>
  seed = 1234,
  level_names = c("sim"),
  n_{per_level} = c(1),
  index_level = FALSE
)
seeds_list12 <- seed_maker(</pre>
  seed = 1234,
  level_names = list("sim", c("component_a", "component_b", "component_c")),
 n_{per_level} = c(1, 3),
  index_level = c(FALSE, TRUE)
)
seeds_list13 <- seed_maker(</pre>
  seed = 1234,
  level_names = list("sim", c("component_a", "component_b", "component_c")),
  n_{per_level} = c(1, 3),
  index_level = c(FALSE, FALSE)
)
identical(seeds_list12, seeds_list13)
seeds_flat1 <- seed_maker(</pre>
  seed = 1234,
  level_names = c("sim"),
  n_{per_level} = c(10),
  flatten = TRUE
)
is.list(seeds_flat1)
is.integer(seeds_flat1)
seeds_flat2 <- seed_maker(</pre>
  seed = 1234,
  level_names = c("sim", "dataset"),
  n_{per_level} = c(10, 15),
  flatten = TRUE
)
is.list(seeds_flat2$sim1)
is.integer(seeds_flat2$sim1)
seeds_flat5 <- seed_maker(</pre>
  seed = 1234,
  level_names = list(
    "sim",
    c("component_a", "component_b", "component_c"),
    "dataset"
```

```
),
  n_{per_level} = c(10, 3, 5),
  flatten = TRUE
)
is.list(seeds_flat5$sim1$component_a)
is.integer(seeds_flat5$sim1$component_a)
seeds_flat9 <- seed_maker(</pre>
  seed = 1234,
  level_names = c("sim"),
  n_{per_level} = c(1),
  flatten = TRUE
is.list(seeds_flat9)
is.integer(seeds_flat9)
try(seeds_flat10 <- seed_maker(</pre>
  seed = 1234,
  level_names = c("sim"),
  n_per_level = c(10),
  index_level = FALSE,
  flatten = TRUE
))
seeds_flat11 <- seed_maker(</pre>
  seed = 1234,
  level_names = c("sim"),
  n_per_level = c(1),
  index_level = FALSE,
  flatten = TRUE
)
is.list(seeds_flat11)
is.integer(seeds_flat11)
```

## **Index**

```
seed_gen, 2
seed_maker, 3
set.seed(), 2, 3
```