

# Package ‘RcppMovStat’

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**Author** Pengfei (Leonard) Li

**Maintainer** Pengfei (Leonard) Li <pli3@tulane.com>

**Description** This is a package providing several efficient functions to calculate common moving(or rolling, running) statitics for both EVENLY and UN-EVENLY SPACED Time Series: moving average, moving median, and moving maximum(minimum) . Built on C++, these functions would be apparently more efficient than those written in traditional R and also faster than Others written by Rcpp.

**Suggests** zoo, caTools, Rcpp

**License** GPL (>= 2)

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RcppMovStat-package      *Fast Moing Statitics Caculation via Rcpp*


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### Description

This is a package providing several efficient functions to calculate common moving(or rolling, running) statistics: moving average, moving median, and moving maximum(minimum) . Built on C++, these functions are apparently more efficient than those written in traditional R.

### Details

See the following instructions for further details.

### Author(s)

Pengfei (Leonard) Li, pli3@tulane.edu.

Maintainer: Pengfei (Leonard) Li <pli3@tulane.edu>

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movCount      *Moving Count*


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### Description

This function returns a moving count of the given vector.

### Usage

```
movCount(vec, n = 1L, ss = 1L, na_rm = FALSE, sizeD = FALSE,
         align = "left")
```

```
movCountr(vec, n = 1L, ss = 1L, na_rm = FALSE, sizeD = FALSE)
```

### Arguments

vec	A numeric vector.
n	An integer: moving window size, with 1 as default
ss	An integer: step size, only calculating at points with an equal distance <i>ss</i> . Namely, there are <i>ss-1</i> number between each two 'consecutive' points
na_rm	logical. Should missing values (including NaN) be removed?
sizeD	logical. Only applied when <i>ss &gt; 1</i> , it decides whether to get a result of smaller size. If <i>sizeD = T</i> , <i>align</i> does not affect the output.
align	A string denotes how to align the moving average, three options: "left", "middle", "right"

### Details

This function counts the number of non-missing values for each moving window. It is especially designed for *vec* vector with missing values. Otherwise, it will return a trival vector with all elements *n*.

**Value**

This function returns a vector whose length is the same as that of *vec* or is `ceiling(( $L - n + 1$ )/ss)`, (when *sizeD* = *T*), where *L* is the length of *vec*.

**Functions**

- `movCountr`: An function equivalent to `movCount(..., align = "right")`

**Examples**

```
movCount(c(1, 4, 3, NA, 8), 2, na_rm = TRUE)
movCount(c(1, 4, 3, NA, 8), 2, na_rm = TRUE, align = 'right')
movCountr(c(1, 4, 3, NA, 8), 2, na_rm = TRUE)
movCount(c(1, 4, 3, NA, NA), 2, na_rm = TRUE)
```

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movCountUE

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*Weighted Simple Moving Count for Unevenly Spaced Time Series*


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**Description**

This function returns A matrix: the first column is the position, the second column the input vector, and third column Moving Count of the given vector. The weight argument is optional.

**Usage**

```
movCountUE(vec, pos, n = 1L, ss = 1L, na_rm = FALSE, sizeD = FALSE,
  align = "left")
```

```
movCountUEr(vec, pos, n = 1L, ss = 1L, na_rm = FALSE, sizeD = FALSE)
```

**Arguments**

<code>vec</code>	A numeric vector.
<code>pos</code>	A numeric vector with all integers. Its length must be the SAME as <i>vec</i> . N.B. We use integers to represent the (relative) positions of every point. The first element MUST BE 1, which is design to follow THE 1-INDEXED RULE of R.
<code>n</code>	An integer: moving window size, with 1 as default
<code>ss</code>	An integer: step size, only calculating at points with an equal distance <i>ss</i> . Namely, there are <i>ss</i> -1 number between each two 'consecutive' points
<code>na_rm</code>	logical. Should missing values (including NaN) be removed?
<code>sizeD</code>	logical. Only applied when <i>ss</i> > 1, it decides whether to get a result of smaller size. If <i>sizeD</i> = <i>T</i> , <i>align</i> does not affect the output.
<code>align</code>	A string denotes how to align the moving average, three options: "left", "middle", "right"

## Details

This function counts the number of non-missing values for each moving window. It is especially designed for *vec* vector with missing values. Otherwise, it will return a trivial vector with all elements  $n$ .

This function is more helpful than `movCount`, as we would have missing values for an Unevenly Spaced Time Series.

For matrix details, please refer to details of `movMeanUE`.

## Value

This function returns A MATRIX of size:  $L * 3$ , where  $L$  is the length of vector, or of size:  $L1 * 3$ , where  $L1 = \text{ceiling}((nrow - n + 1)/ss)$ , (when  $sizeD = T$ ). In the matrix, the first column denotes the position, the second column the original vector, and the third column the moving average.

## Functions

- `movCountUEr`: An function equivalent to `movCountUE(..., align = "right")`

## Examples

```
movCountUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), 2)
movCountUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, na_rm = TRUE)
movCountUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3, na_rm = TRUE,
sizeD = TRUE)
movCountUE(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2,
ss = 3, na_rm = TRUE, align = "right")
movCountUE(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3,
na_rm = TRUE, sizeD = TRUE, align = "right")
movCountUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3,
na_rm = TRUE)
movCountUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3, na_rm = TRUE,
sizeD = TRUE)
```

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movEmean

*Basic Exponential Moving Mean*

---

## Description

This function returns a basic exponential moving average (EMA) of the given vector.

## Usage

```
movEmean(vec, n = 1L, smFac = NULL)
```

## Arguments

<code>vec</code>	A numeric vector.
<code>n</code>	An integer: moving window size, with 1 as default
<code>smFac</code>	A number: smoothing factor, with default $2/(n + 1)$ , see <b>details</b> below.

## Details

This function makes fairly efficient the computation of EMA, which dubbed as basic exponential smoothing, the same section of [https://en.wikipedia.org/wiki/Exponential\\_smoothing](https://en.wikipedia.org/wiki/Exponential_smoothing). It provides an access to define *smFac* yourself, i.e the smoothing factor, whose default is  $2/(n + 1)$ .

## Value

This function returns a vector whose length is the same as that of *vec*.

## Examples

```
movMean(c(1, 4, 3, 6, 8), 2, smFac = 1/3)
movMean(c(1, 4, 3, 6, 8), 2)
```

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movMean	<i>Weighted Simple Moving Mean</i>
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## Description

This function returns a simple moving average of the given vector. The weight argument is optional.

## Usage

```
movMean(vec, n = 1L, ss = 1L, w = NULL, na_rm = FALSE, sizeD = FALSE,
        align = "left")

movMeanr(vec, n = 1L, ss = 1L, w = NULL, na_rm = FALSE, sizeD = FALSE)
```

## Arguments

<i>vec</i>	A numeric vector.
<i>n</i>	An integer: moving window size, with 1 as default
<i>ss</i>	An integer: step size, only calculating at points with an equal distance <i>ss</i> . Namely, there are <i>ss-1</i> number between each two 'consecutive' points
<i>w</i>	An optional weight vector of length <i>n</i> . It will be automatically normalized (sum to 1).
<i>na_rm</i>	logical. Should missing values (including NaN) be removed?
<i>sizeD</i>	logical. Only applied when <i>ss &gt; 1</i> , it decides whether to get a result of smaller size. If <i>sizeD = T</i> , <i>align</i> does not affect the output.
<i>align</i>	A string denotes how to align the moving average, three options: "left", "middle", "right"

## Details

Despite of Efficient computation, usually 5~6 times faster than the moving average function in package [roll\\_mean](#), it is able to handle potential missing values (*NA* or *NaN*) in the *vec*. For instace, the output of the second examle is *NA, NA, 2.2000003.7142864.875000*. The last number 5.5 is obtained by using renormalized weight, namely omitting 0.2. The weight applied would be  $0.5/(0.5 + 0.3)$  and  $0.3/(0.5 + 0.3)$ . Hence,

$$4.875 = 3 * 0.5/(0.5 + 0.3) + 8 * 0.3/(0.5 + 0.3)$$

**Value**

This function returns a vector whose length is the same as that of *vec* or is `ceiling(( $L - n + 1$ )/ss)`, (when *sizeD* = *T*), where *L* is the length of *vec*.

**Functions**

- `movMeanr`: An function equivalent to `movMean(..., align = "right")`

**Examples**

```
movMean(c(1, 4, 3, NA, 8), 3, align = "right", na_rm = TRUE)
movMean(c(1, 4, 3, NA, 8), 3, w = c(0.5, 0.2, 0.3), na_rm = TRUE, align = "right")
movMean(c(1, 4, 3, NA, 8, 4, 5, 9, 6, 0), n = 3, ss = 4, na_rm = TRUE, align = "right")
```

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movMeanUE	<i>Weighted Simple Moving Mean for Unevenly Spaced Time Series</i>
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---

**Description**

This function returns A matrix: the first column is the position, the second column the input vector, and third column simple moving average of the given vector. The weight argument is optional.

**Usage**

```
movMeanUE(vec, pos, n = 1L, ss = 1L, w = NULL, na_rm = FALSE,
  sizeD = FALSE, align = "left")

movMeanUEr(vec, pos, n = 1L, ss = 1L, w = NULL, na_rm = FALSE,
  sizeD = FALSE)
```

**Arguments**

<code>vec</code>	A numeric vector.
<code>pos</code>	A numeric vector with all integers. Its length must be the SAME as <i>vec</i> . N.B. We use integers to represent the (relative) positions of every point. The first element MUST BE 1, which is design to follow THE 1-INDEXED RULE of R.
<code>n</code>	An integer: moving window size, with 1 as default
<code>ss</code>	An integer: step size, only calculating at points with an equal distance <i>ss</i> . Namely, there are <i>ss-1</i> number between each two 'consecutive' points
<code>w</code>	An optional weight vector of length <i>n</i> . It will be automatically normalized (sum to 1).
<code>na_rm</code>	logical. Should missing values (including NaN) be removed?
<code>sizeD</code>	logical. Only applied when <i>ss</i> > 1, it decides whether to get a result of smaller size. If <i>sizeD</i> = <i>T</i> , <i>align</i> does not affect the output.
<code>align</code>	A string denotes how to align the moving average, three options: "left", "middle", "right"

## Details

This function is especially designed for Unevenly Spaced Time Series. It is efficient as it inherits the similar routine of `movMean`.

The result is kind of tricky. To make it clear, it is written to return a MATRIX. For instance, the third column of the output of second example is `NA, 2.5, 4.0, NA, NA, NA, 3.0, 3.0, 8.0`. 2.5 is the average of 1 and 4, and 4.0 the average of 4. The third column of the output of third example is the every third element starting from  $n$ th number.

For how weights,  $w$ , work, one can refer to `movMean`.

## Value

This function returns A MATRIX of size:  $L * 3$ , where  $L$  is the length of vector, or of size:  $L1 * 3$ , where  $L1 = \text{ceiling}((nrow - n + 1)/ss)$ , (when  $sizeD = T$ ). In the matrix, the first column denotes the position, the second column the original vector, and the third column the moving average.

## Functions

- `movMeanUEr`: An function equivalent to `movMeanUE(..., align = "right")`

## Examples

```
movMeanUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), 2)
movMeanUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, na_rm = TRUE)
movMeanUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3, na_rm = TRUE,
sizeD = TRUE)
movMeanUE(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), w = c(0, 1), n = 2,
ss = 3, na_rm = TRUE, align = "right")
movMeanUE(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3,
na_rm = TRUE, sizeD = TRUE, align = "right")
movMeanUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3,
na_rm = TRUE)
movMeanUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3, na_rm = TRUE,
sizeD = TRUE)
```

---

movQt

*Moving Quantile(Moving Median, Moving Minimum, Moving Maximum)*

---

## Description

This function returns a moving quantile of the given vector.

## Usage

```
movQt(vec, n = 1L, prob = 0.5, ss = 1L, na_rm = FALSE, sizeD = FALSE,
align = "left")

movQtr(vec, n = 1L, prob = 0.5, ss = 1L, na_rm = FALSE, sizeD = FALSE)
```

## Arguments

<code>vec</code>	A numeric vector.
<code>n</code>	An integer: moving window size, with 1 as default
<code>prob</code>	A number: between 0 and 1, meaning <i>prob</i> quantile
<code>ss</code>	An integer: step size, only calculating at points with an equal distance <i>ss</i> . Namely, there are <i>ss</i> -1 number between each two 'consecutive' points
<code>na_rm</code>	logical. Should missing values (including NaN) be removed?
<code>sizeD</code>	logical. Only applied when <i>ss</i> > 1, it decides whether to get a result of smaller size. If <i>sizeD</i> = T, <i>align</i> does not affect the output.
<code>align</code>	A string denotes how to align the moving average, three options: "left", "middle", "right"

## Details

Despite of Efficient computation, this function can return differents kinds of moving quantile, e.g. moving median(*prob* = 0.5), moving minimum(*prob* = 0), and moving maximum(*prob* = 1). It can handle potential missing values(NA or NaN) in the *vec*. When we move to one specific fragment, missing values can be removed by setting *na\_rm* = TRUE. If all values of this fragment is missing, it will return NA.

In terms of the quantile algorithm, please consult type 7 in function [quantile](#).

## Value

This function returns a vector whose length is the same as that of *vec* or is  $\text{ceiling}((L-n+1)/ss)$ , (when *sizeD* = T), where *L* is the length of *vec*.

## Functions

- `movQtr`: An function equivalent to `movQt(..., align = "right")`

## Examples

```
movQt(vec = c(1, 4, 3, NA, 8, 4, 5, 9, 6, 0), n = 3, ss = 4, na_rm = TRUE, align = "right")
movQt(vec = c(1, 4, 3, NA, 8, 4, 5, 9, 6, 0), n = 3, na_rm = TRUE, align = "right")
movQt(vec = c(1, 4, 3, NA, NA, NA, 5, 9, 6, 0), n = 3, ss = 4, na_rm = TRUE, align = "middle")
```

---

<code>movQtUE</code>	<i>Moving quantile_UE(Moving Median, Moving Minimum, Moving Maximum) for Unevenly Spaced Time Series</i>
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---

## Description

This function returns A matrix: the first column is the position, the second column the input vector, and third column moving quantile\_UE of the given vector.

## Usage

```
movQtUE(vec, pos, n = 1L, prob = 0.5, ss = 1L, na_rm = FALSE,
        sizeD = FALSE, align = "left")

movQtUEr(vec, pos, n = 1L, prob = 0.5, ss = 1L, na_rm = FALSE,
        sizeD = FALSE)
```



## Arguments

<code>vec</code>	A numeric vector.
<code>pos</code>	A numeric vector with all integers. Its length must be the SAME as <i>vec</i> . N.B. We use integers to represent the (relative) positions of every point. The first element MUST BE 1, which is design to follow THE 1-INDEXED RULE of R.
<code>n</code>	An integer: moving window size, with 1 as default
<code>prob</code>	A number: between 0 and 1, meaning <i>prob</i> quantile_UE
<code>ss</code>	An integer: step size, only calculating at points with an equal distance <i>ss</i> . Namely, there are <i>ss-1</i> number between each two 'consecutive' points
<code>na_rm</code>	logical. Should missing values (including NaN) be removed?
<code>sizeD</code>	logical. Only applied when <i>ss</i> > 1, it decides whether to get a result of smaller size. If <i>sizeD</i> = <i>T</i> , <i>align</i> does not affect the output.
<code>align</code>	A string denotes how to align the moving average, three options: "left", "middle", "right"

## Details

This function is especially designed for Unevenly Spaced Time Series. It is efficient as it herits the similiar routine of `movQt`.

The result is kind of tricky. To make it clear, it is written to return a MATRIX. For instance, the third column of the output of second example is 2.5, *NA*, *NA*, *NA*, *NA*, *NA*, 3.0, *NA*, *NA*. 2.5 is the median of 1 and 4, and 4.0 the average of 4. The third column of the output of third example is the every third element starting from *n*th number.

For how weights, *w*, work, one can refer to `movQt`.

## Value

This function returns A MATRIX of size:  $L * 3$ , where L is the length of vector, or of size:  $L1 * 3$ , where  $L1 = \text{ceiling}((nrow - n + 1)/ss)$ , (when *sizeD* = *T*). In the matrix, the first column denotes the position, the second column the original vector, and the third column the moving average.

## Functions

- `movQtUER`: An function equivalent to `movQtUE(..., align = "right")`

## Examples

```
movQtUE(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3, na_rm = TRUE)
movQtUE(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3, na_rm = TRUE, sizeD = TRUE)
movQtUE(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, na_rm = TRUE, align = "middle")
movQtUE(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3, na_rm = TRUE, sizeD = TRUE,
  align = "right")
```

---

movSum	<i>Weighted Simple Moving Sum</i>
--------	-----------------------------------

---

## Description

This function returns a simple moving sum of the given vector. The weight argument is optional.

## Usage

```
movSum(vec, n = 1L, ss = 1L, w = NULL, na_rm = FALSE, sizeD = FALSE,
       align = "left")
```

```
movSumr(vec, n = 1L, ss = 1L, w = NULL, na_rm = FALSE, sizeD = FALSE)
```

## Arguments

vec	A numeric vector.
n	An integer: moving window size, with 1 as default
ss	An integer: step size, only calculating at points with an equal distance <i>ss</i> . Namely, there are <i>ss-1</i> number between each two 'consecutive' points
w	An optional weight vector of length <i>n</i> .
na_rm	logical. Should missing values (including NaN) be removed?
sizeD	logical. Only applied when <i>ss</i> > 1, it decides whether to get a result of smaller size. If <i>sizeD</i> = <i>T</i> , <i>align</i> does not affect the output.
align	A string denotes how to align the moving average, three options: "left", "middle", "right"

## Details

This function can obtain the moving sum efficiently. It serves as somehow a generalized version of [movMean](#). The difference is that it will not automatically normalized the weights vector, *w* argument. If there is no missing value in *vec*, and *w* is normalized, which means the sum of all elements is 1, this function will return a moving average.

## Value

This function returns a vector whose length is the same as that of *vec* or is `ceiling((L-n+1)/ss)`, (when *sizeD* = *T*), where *L* is the length of *vec*.

## Functions

- `movSumr`: An function equivalent to `movMean(..., align = "right")`

## Examples

```
movSum(c(1, 4, 3, NA, 8), 3, align = "right", na_rm = TRUE)
movSum(c(1, 4, 3, NA, 8), 3, w = c(0.5, 0.2, 0.3), na_rm = TRUE, align = "right")
movSum(c(1, 4, 3, NA, 8, 4, 5, 9, 6, 0), n = 3, ss = 4, na_rm = TRUE, align = "right")
```

---

movSumUE

---

Weighted Simple Moving Sum for Unevenly Spaced Time Series

---

## Description

This function returns A matrix: the first column is the position, the second column the input vector, and third column moving sum of the given vector. The weight argument is optional.

## Usage

```
movSumUE(vec, pos, n = 1L, ss = 1L, w = NULL, na_rm = FALSE,
         sized = FALSE, align = "left")
```

```
movSumUEr(vec, pos, n = 1L, ss = 1L, w = NULL, na_rm = FALSE,
          sized = FALSE)
```

## Arguments

vec	A numeric vector.
pos	A numeric vector with all integers. Its length must be the SAME as <i>vec</i> . N.B. We use integers to represent the (relative) positions of every point. The first element MUST BE 1, which is design to follow THE 1-INDEXED RULE of R.
n	An integer: moving window size, with 1 as default
ss	An integer: step size, only calculating at points with an equal distance <i>ss</i> . Namely, there are <i>ss-1</i> number between each two 'consecutive' points
w	An optional weight vector of length <i>n</i> . It will be automatically normalized (sum to 1).
na_rm	logical. Should missing values (including NaN) be removed?
sized	logical. Only applied when <i>ss</i> > 1, it decides whether to get a result of smaller size. If <i>sizeD</i> = T, <i>align</i> does not affect the output.
align	A string denotes how to align the moving average, three options: "left", "middle", "right"

## Details

This function can obtain the moving sum efficiently. It serves as somehow a generalized version of [movMeanUE](#). The difference is that it will not automatically normalized the weights vector, *w* argument.

If there is no missing value in *vec*, and *w* is normalized, which means the sum of all elements is 1, this function will return a moving average. For matrix details, please refer to details of [movMeanUE](#).

## Value

This function returns A MATRIX of size:  $L * 3$ , where *L* is the length of vector, or of size:  $L1 * 3$ , where  $L1 = \text{ceiling}((nrow - n + 1)/ss)$ , (when *sizeD* = T). In the matrix, the first column denotes the position, the second column the original vector, and the third column the moving average.

**Functions**

- movSumUEr: An function equivalent to movSumUE(..., align = "right")

**Examples**

```
movSumUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), 2)
movSumUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, na_rm = TRUE)
movSumUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3, na_rm = TRUE,
sizeD = TRUE)
movSumUE(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), w = c(0, 1), n = 2,
ss = 3, na_rm = TRUE, align = "right")
movSumUE(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3,
na_rm = TRUE, sizeD = TRUE, align = "right")
movSumUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3,
na_rm = TRUE)
movSumUEr(c(1, 4, 3, NA, 8), pos = c(1, 2, 7, 8, 9), n = 2, ss = 3, na_rm = TRUE,
sizeD = TRUE)
```

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