

Package ‘fiber’

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Type Package

Title S7 Data Structures for Diffusion MRI Tractography

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Description Provides three S7 classes — streamline, bundle, and bundle_set — for representing diffusion MRI tractography data in R, together with a concise set of methods for computing shape descriptors (arc-length, curvature, torsion, sinuosity), the Hausdorff distance between streamlines, arc-length reparametrization of streamlines and bundles onto uniform grids, combination of streamlines or bundles into a single bundle, combination of bundles from multiple subjects or sessions into a bundle_set, and coercion to and from the dwiFiber S4 class of the 'dti' package. See Dell'Acqua, F., Descoteaux, M. and Leemans, A. (2024) ``Handbook of Diffusion MR Tractography" <[doi:10.1016/C2018-0-02520-7](https://doi.org/10.1016/C2018-0-02520-7)> for more about the mathematical and computational underpinnings of diffusion MRI tractography.

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Author Aymeric Stamm [aut, cre] (ORCID:
<<https://orcid.org/0000-0002-8725-3654>>)

Maintainer Aymeric Stamm <aymeric.stamm@cnrs.fr>

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add_shape_descriptors *Adds shape descriptors to a streamline or bundle*

Description

add_shape_descriptors() is an S7 generic that computes a number of shape descriptors for each [streamline](#) object and stores them in the @streamline_data or @point_data slots as appropriate, with methods available for the following classes:

- [fiber::bundle](#)
- [fiber::streamline](#)

This function provides a convenient way to compute shape descriptors and attach them to [streamline](#) or [bundle](#) objects. See the documentation for each individual shape descriptor function (e.g. [get_euclidean_length\(\)](#), [get_curvilinear_length\(\)](#), [get_sinusosity\(\)](#), [get_curvature\(\)](#), [get_torsion\(\)](#)) for more details on how each descriptor is computed.

For [bundle](#) objects, scalar descriptors (euclidean_length, curvilinear_length, sinusosity) are stored as length-S vectors in bundle@streamline_data. Per-point descriptors (curvature, torsion) continue to be stored in each individual streamline's @point_data. Both are accessible via bundle[[i]]@streamline_data and bundle[[i]]@point_data respectively, through the subsetting push-down mechanism.

Usage

```
add_shape_descriptors(  
  x,  
  descriptors = c("euclidean_length", "curvilinear_length", "sinusosity", "curvature",  
                 "torsion")  
)
```

Arguments

x A [streamline](#) or [bundle](#) object.

descriptors A character vector of shape descriptors to add. Defaults to all available descriptors: c("euclidean_length", "curvilinear_length", "sinusosity", "curvature", "torsion").

Value

An object of the same class as x with the specified shape descriptors added to the appropriate slots.

Examples

```
# add multiple shape descriptors to a single streamline  
sl <- streamline(points = cbind(X = runif(10), Y = runif(10), Z = runif(10)))  
sl <- add_shape_descriptors(  
  sl,
```

```

  descriptors = c("euclidean_length", "curvilinear_length", "sinuosity")
)
sl@streamline_data$euclidean_length

# add multiple shape descriptors to a bundle
sl2 <- streamline(points = cbind(X = runif(20), Y = runif(20), Z = runif(20)))
b <- bundle(streamlines = list(sl, sl2))
b <- add_shape_descriptors(
  b,
  descriptors = c("euclidean_length", "curvilinear_length", "sinuosity")
)
b@streamline_data$euclidean_length # length-2 vector

```

add_shape_descriptors-fiber-bundle-method

[add_shape_descriptors\(\)](#) *method for bundle objects*

Description

Adds shape descriptors to a [bundle](#). Scalar descriptors (`euclidean_length`, `curvilinear_length`, `sinuosity`) are stored as length- S vectors in `bundle@streamline_data`. Per-point descriptors (`curvature`, `torsion`) are stored in each individual streamline's `@point_data`. Both are accessible via the subsetting push-down (`bundle[[i]]`).

Arguments

<code>x</code>	A bundle object.
<code>descriptors</code>	A character vector of shape descriptors to add. Defaults to all available descriptors: <code>c("euclidean_length", "curvilinear_length", "sinuosity", "curvature", "torsion")</code> .

Value

A [bundle](#) with the specified shape descriptors added.

See Also

[add_shape_descriptors\(\)](#)

add_shape_descriptors-fiber-streamline-method

[add_shape_descriptors\(\)](#) *method for streamline objects*

Description

Adds multiple shape descriptors to a single [streamline](#) object.

Arguments

x A [streamline](#) object.

descriptors A character vector of shape descriptors to add. Defaults to all available descriptors: `c("euclidean_length", "curvilinear_length", "sinuosity", "curvature", "torsion")`.

Value

A [streamline](#) with the specified shape descriptors added to the `@streamline_data` or `@point_data` slots as appropriate.

See Also

[add_shape_descriptors\(\)](#)

as_bundle

Coerce an object to a bundle

Description

`as_bundle()` converts a supported object into a [bundle](#).

Usage

```
as_bundle(x, ...)
```

Arguments

x An object to coerce.

... Additional arguments (currently unused).

Details

Currently supported input classes:

- **streamline**: wrapped in a single-element **bundle** (lossless).
- **bundle**: returned unchanged.
- **dwiFiber** (from **dti**): each fiber becomes a **streamline**. The per-point direction vectors (columns 4–6 of `@fibers`) are stored as `@point_data$direction_x`, `@point_data$direction_y`, and `@point_data$direction_z`. Tracking metadata (`method`, `minfa`, `maxangle`) are stored as scalars in `@bundle_data`. Vector metadata (`ddim`, `ddim0`, `voxelext`, `orientation`) are stored as individual scalar entries (e.g. `voxelext_1`, `voxelext_2`, `voxelext_3`).

Value

A **bundle** object.

See Also

[as_streamline\(\)](#), [as_dwifiber\(\)](#)

Examples

```
s1 <- streamline(points = cbind(X = runif(5), Y = runif(5), Z = runif(5)))
b <- as_bundle(s1)
b@n_streamlines # 1
```

as_bundle-fiber-bundle-method

[as_bundle\(\)](#) *method for bundle objects*

Description

[as_bundle\(\)](#) method for bundle objects

Arguments

`x` A **bundle** object.
`...` Additional arguments (currently unused).

Value

`x` unchanged.

See Also

[as_bundle\(\)](#)

as_bundle-fiber-streamline-method

[as_bundle\(\)](#) *method for streamline objects*

Description

[as_bundle\(\)](#) method for streamline objects

Arguments

x A [streamline](#) object.
... Additional arguments (currently unused).

Value

A [bundle](#) containing x as its sole streamline.

See Also

[as_bundle\(\)](#)

as_bundle_set

Coerce an object to a bundle_set

Description

[as_bundle_set\(\)](#) converts a supported object into a [bundle_set](#).

Usage

```
as_bundle_set(x, ...)
```

Arguments

x An object to coerce.
... Additional arguments passed to methods (e.g. name for bundles).

Details

Currently supported input classes:

- [bundle_set](#): returned unchanged.
- [bundle](#): wrapped in a single-element [bundle_set](#). An optional name argument sets the element name (defaults to NULL, producing an unnamed single-element set).

Value

A `bundle_set` object.

See Also

`bundle_set()`, `bind_bundle_sets()`

Examples

```
s1 <- streamline(points = cbind(X = runif(5), Y = runif(5), Z = runif(5)))
b <- bundle(streamlines = list(s1))
bs <- as_bundle_set(b, name = "sub-01")
bs@n_bundles # 1
```

as_dwifiber

Coerce a streamline or bundle to a dwiFiber object

Description

`as_dwifiber()` converts a `streamline` or `bundle` to the S4 class `dwiFiber` from the **dti** package.

Usage

```
as_dwifiber(x, ...)
```

Arguments

`x` A `streamline` or `bundle` object.
`...` Additional arguments (currently unused).

Details

Per-point direction vectors are taken from `@point_data$direction_x`, `@point_data$direction_y`, and `@point_data$direction_z` when present; otherwise they are estimated via finite differences of the coordinates (forward difference at the first point, backward difference at the last, central differences in between), then unit-normalised.

Bundle-level metadata stored in `@bundle_data` under the keys `method`, `minfa`, `maxangle`, `level`, and `source` are transferred to the corresponding `dwiFiber` slots when present. Vector-valued fields such as `ddim`, `ddim0`, `voxelx`, and `orientation` must be stored as individual scalars (e.g. `ddim_1`, `ddim_2`, `ddim_3`) and are reconstructed into vectors for the `dwiFiber` object.

Value

An S4 object of class `dwiFiber` (from **dti**).

See Also

`as_streamline()`, `as_bundle()`

Examples

```
if (requireNamespace("dti", quietly = TRUE)) {  
  sl <- streamline(points = cbind(X = runif(5), Y = runif(5), Z = runif(5)))  
  b <- bundle(streamlines = list(sl))  
  dfi <- as_dwifiber(b)  
  class(dfi) # "dwiFiber"  
}
```

as_dwifiber-fiber-bundle-method

[as_dwifiber\(\)](#) *method for bundle objects*

Description

[as_dwifiber\(\)](#) method for bundle objects

Arguments

x A **bundle** object.
... Additional arguments (currently unused).

Value

An S4 dwiFiber object.

See Also

[as_dwifiber\(\)](#)

as_dwifiber-fiber-streamline-method

[as_dwifiber\(\)](#) *method for streamline objects*

Description

[as_dwifiber\(\)](#) method for streamline objects

Arguments

x A **streamline** object.
... Additional arguments (currently unused).

Value

An S4 dwiFiber object.

See Also

[as_dwifiber\(\)](#)

as_streamline *Coerce an object to a streamline*

Description

as_streamline() converts a supported object into a [streamline](#).

Usage

```
as_streamline(x, ...)
```

Arguments

x	An object to coerce.
...	Additional arguments (currently unused).

Details

Currently supported input classes:

- [streamline](#): returned unchanged.
- `dwiFiber` (from `dti`): the object must contain **exactly one** fiber. For multi-fiber objects use [as_bundle\(\)](#) instead.

Value

A [streamline](#) object.

See Also

[as_bundle\(\)](#), [as_dwifiber\(\)](#)

Examples

```
s1 <- streamline(points = cbind(X = runif(5), Y = runif(5), Z = runif(5)))
identical(as_streamline(s1), s1) # TRUE - identity coercion
```

as_streamline-fiber-bundle-method

[as_streamline\(\)](#) *method for bundle objects*

Description

[as_streamline\(\)](#) method for bundle objects

Arguments

x A [bundle](#) object containing exactly one streamline.
... Additional arguments (currently unused).

Value

The sole [streamline](#) inside x.

See Also

[as_streamline\(\)](#)

as_streamline-fiber-streamline-method

[as_streamline\(\)](#) *method for streamline objects*

Description

[as_streamline\(\)](#) method for streamline objects

Arguments

x A [streamline](#) object.
... Additional arguments (currently unused).

Value

x unchanged.

See Also

[as_streamline\(\)](#)

bind_bundles	<i>Combine streamlines and/or bundles into a single bundle</i>
--------------	--

Description

Accepts any mix of [streamline](#) and [bundle](#) objects. All streamlines are collected into a flat list and wrapped in a new [bundle](#). `bundle_data` from the first [bundle](#) argument (if any) is preserved; pass your own via the `bundle_data` argument to override. Similarly for `streamline_data`.

Usage

```
bind_bundles(..., streamline_data = NULL, bundle_data = NULL)
```

Arguments

<code>...</code>	One or more streamline or bundle objects.
<code>streamline_data</code>	A named list of per-streamline vectors to attach to the resulting bundle . Defaults to the <code>@streamline_data</code> of the first bundle input if one is present.
<code>bundle_data</code>	A named list of bundle-level scalar metadata to attach to the resulting bundle . Defaults to an empty list (or the <code>bundle_data</code> of the first bundle input if one is present and <code>bundle_data</code> is not supplied).

Value

A [bundle](#) containing all input streamlines.

Examples

```
s11 <- streamline(points = cbind(X = runif(5), Y = runif(5), Z = runif(5)))
s12 <- streamline(points = cbind(X = runif(5), Y = runif(5), Z = runif(5)))
b1 <- bundle(streamlines = list(s11))
b2 <- bundle(streamlines = list(s12))

# combine two bundles
b_all <- bind_bundles(b1, b2)
b_all@n_streamlines # 2

# mix a bundle and a loose streamline
b_mixed <- bind_bundles(b1, s12)
b_mixed@n_streamlines # 2
```

bind_bundle_sets	<i>Combine bundles and/or bundle_sets into a single bundle_set</i>
------------------	--

Description

Accepts any mix of `bundle` objects or `bundle_set` objects. All bundles are collected into a flat list and wrapped in a new `bundle_set`. Bare `bundle` arguments may optionally be named; unnamed bundles are included without a name label.

Usage

```
bind_bundle_sets(..., bundle_data = NULL, set_data = NULL)
```

Arguments

...	<code>bundle</code> objects or <code>bundle_set</code> objects to combine. Named bare <code>bundle</code> arguments will carry their name into the resulting set.
bundle_data	A named list of per-bundle vectors to attach to the resulting <code>bundle_set</code> . Defaults to the <code>@bundle_data</code> of the first <code>bundle_set</code> input if present.
set_data	A named list of set-level scalar metadata to attach to the resulting <code>bundle_set</code> . Defaults to the <code>set_data</code> of the first <code>bundle_set</code> input (if present) or an empty list.

Value

A `bundle_set` containing all input bundles.

Examples

```
s1 <- streamline(points = cbind(X = runif(5), Y = runif(5), Z = runif(5)))
b1 <- bundle(streamlines = list(s1))
b2 <- bundle(streamlines = list(s1))

# two named bare bundles
bs <- bind_bundle_sets("sub-01" = b1, "sub-02" = b2)
bs@n_bundles # 2
bs@bundle_data$if_from_input_list # c("sub-01", "sub-02")

# combine two bundle_sets
bs1 <- bundle_set(list("sub-01" = b1))
bs2 <- bundle_set(list("sub-02" = b2))
bs_all <- bind_bundle_sets(bs1, bs2)
bs_all@n_bundles # 2
```

bundle

*Bundle S7 class***Description**

A bundle is an ordered collection of [streamline](#) objects representing a tractogram or white-matter bundle. It stores three compartments:

- `@streamlines` — a list of [streamline](#) objects.
- `@streamline_data` — a named list of per-streamline vectors (any type), each of length S (the number of streamlines). Values common to all streamlines may be lifted here automatically from the individual streamlines' `@streamline_data` slots at construction time.
- `@bundle_data` — a named list of scalars (length-1 values, any type) holding bundle-level metadata.

Usage

```
bundle(streamlines = list(), streamline_data = list(), bundle_data = list())
```

Arguments

`streamlines` A list of [streamline](#) objects.

`streamline_data` A named list of per-streamline vectors of length S . If not supplied, any `@streamline_data` keys common to **all** streamlines are lifted automatically.

`bundle_data` A named list of bundle-level scalar metadata.

Value

A bundle S7 object.

Methods for standard generics

The following methods are defined for bundle objects:

- `format(x, ...)`: Returns a cli-formatted string describing the bundle object.
- `print(x, ...)`: Prints the formatted string to the console and invisibly returns `x`.
- `length(x)`: Returns the number of streamlines (equivalent to `x@n_streamlines`).
- `x[[i]]`: Extracts the i -th [streamline](#) from the bundle, with bundle-level `@streamline_data` pushed back into the streamline.
- `x[i]`: Returns a new [bundle](#) containing only the selected streamlines, with `@bundle_data` and the subset of `@streamline_data` preserved.

Additional properties

@n_streamlines An integer scalar giving the number of streamlines in the bundle (read-only).

@streamline_attributes A character vector of the names of the per-streamline attributes stored at the bundle level (read-only).

@bundle_attributes A character vector of the names of the bundle-level attributes (read-only).

Examples

```
s11 <- streamline(
  points = cbind(X = 0:4, Y = 0:4, Z = 0:4),
  streamline_data = list(mean_FA = 0.6, label = "CST")
)
s12 <- streamline(
  points = cbind(X = 1:3, Y = 1:3, Z = 1:3),
  streamline_data = list(mean_FA = 0.7, label = "CST")
)
# mean_FA and label are common to both streamlines and are lifted
b <- bundle(streamlines = list(s11, s12))
b@n_streamlines # 2
b@streamline_attributes # c("mean_FA", "label")
b@streamline_data$mean_FA # c(0.6, 0.7)

# Subsetting pushes streamline_data back down
b[[1]]@streamline_data$mean_FA # 0.6
b[1]@streamline_data$mean_FA # c(0.6)
```

 bundle_set

Bundle set S7 class

Description

A `bundle_set` is a collection of `bundle` objects, designed for multi-subject or multi-session studies where each element represents one subject's (or session's) tractogram. It stores three compartments:

- @bundles — a list of `bundle` objects (names are optional).
- @bundle_data — a named list of per-bundle vectors (any type), each of length B (the number of bundles). Values common to all bundles may be lifted here automatically from the individual bundles' @bundle_data slots at construction time.
- @set_data — a named list of scalars (length-1 values, any type) holding set-level metadata.

Usage

```
bundle_set(bundles = list(), bundle_data = list(), set_data = list())
```

Arguments

<code>bundles</code>	A list of <code>bundle</code> objects (may be named or unnamed).
<code>bundle_data</code>	A named list of per-bundle vectors of length B. If not supplied, any <code>@bundle_data</code> keys common to all bundles are lifted automatically.
<code>set_data</code>	A named list of set-level scalar metadata.

Value

A `bundle_set` S7 object.

Methods for standard generics

The following methods are defined for `bundle_set` objects:

- `format(x, ...)`: Returns a styled character string.
- `print(x, ...)`: Prints the formatted string to the console and invisibly returns `x`.
- `length(x)`: Returns the number of bundles.
- `x[[i]]`: Extracts the *i*-th (or named) `bundle` from the set, with set-level `@bundle_data` pushed back into the bundle.
- `x[i]`: Returns a new `bundle_set` containing only the selected bundles, with `@set_data` and the subset of `@bundle_data` preserved.

Additional properties

`@n_bundles` An integer scalar giving the number of bundles in the set (read-only).

`@bundle_attributes` A character vector of the names of the per-bundle attributes stored at the set level (read-only).

`@set_attributes` A character vector of the names of the set-level attributes (read-only).

Examples

```
s1 <- streamline(points = cbind(X = 1:5, Y = 1:5, Z = 1:5))
b1 <- bundle(
  streamlines = list(s1),
  bundle_data = list(subject = "sub-01")
)
b2 <- bundle(
  streamlines = list(s1),
  bundle_data = list(subject = "sub-02")
)
# subject is common across bundles and lifted to bundle_set@bundle_data
bs <- bundle_set(bundles = list("sub-01" = b1, "sub-02" = b2))
bs@n_bundles           # 2
bs@bundle_attributes   # "subject"
bs@bundle_data$subject # c("sub-01", "sub-02")
```

 compute_hausdorff_distance

Computes the Hausdorff distance between streamlines

Description

compute_hausdorff_distance() is an S7 generic that computes the symmetric Hausdorff distance between [streamline](#) objects based on their 3-D coordinate matrices, with methods available for the following classes:

- ANY
- [fiber::bundle](#)
- [fiber::streamline](#)

The four dispatch cases are:

- [streamline](#) + [streamline](#): returns a single numeric scalar — the symmetric Hausdorff distance between the two streamlines.
- [bundle](#) + **missing**: returns a symmetric numeric distance matrix of dimension $n \times n$, where n is the number of streamlines in the bundle, giving all pairwise Hausdorff distances.
- [bundle](#) + [streamline](#): returns a numeric vector of length n giving the Hausdorff distance from y to each streamline in x .
- [bundle](#) + [bundle](#): returns a symmetric numeric distance matrix of dimension $(n_x + n_y) \times (n_x + n_y)$, treating the concatenation of all streamlines from x and y as one collection.

Usage

```
compute_hausdorff_distance(x, y = NULL)
```

Arguments

- | | |
|----------------|--|
| <code>x</code> | A streamline or bundle object. |
| <code>y</code> | A streamline or bundle object, or NULL (default). When NULL and <code>x</code> is a bundle , the pairwise distance matrix within <code>x</code> is returned. |

Value

- A non-negative numeric scalar when both `x` and `y` are [streamlines](#).
- A [dist](#) object of size n when `x` is a [bundle](#) and `y` is NULL or a [bundle](#) (use `as.matrix()` to expand to a full $n \times n$ matrix).
- A numeric vector of length n when `x` is a [bundle](#) and `y` is a [streamline](#).

Examples

```

s11 <- streamline(points = cbind(X = runif(10), Y = runif(10), Z = runif(10)))
s12 <- streamline(points = cbind(X = runif(10), Y = runif(10), Z = runif(10)))

# streamline x streamline -> scalar
compute_hausdorff_distance(s11, s12)

# bundle x missing -> pairwise dist object
b <- bundle(streamlines = list(s11, s12))
compute_hausdorff_distance(b)
as.matrix(compute_hausdorff_distance(b))

# bundle x streamline -> vector
compute_hausdorff_distance(b, s11)

# bundle x bundle -> combined pairwise matrix
b2 <- bundle(streamlines = list(s12))
compute_hausdorff_distance(b, b2)

```

compute_hausdorff_distance, any-method

[compute_hausdorff_distance\(\)](#) *catch-all method*

Description

[compute_hausdorff_distance\(\)](#) catch-all method

Value

Does not return a value; always throws an error for unsupported input types.

compute_hausdorff_distance-fiber-bundle-method

[compute_hausdorff_distance\(\)](#) *method for bundle objects*

Description

Dispatches to one of three behaviours depending on y:

Arguments

x	A bundle object.
y	NULL, a streamline , or a bundle .

Details

- `y = NULL`: pairwise distances within `x` as a `dist` object of size n , computed in C++ via a single linear loop.
- `y` is a `streamline`: numeric vector of distances from `y` to each streamline in `x`.
- `y` is a `bundle`: `dist` object for the concatenation of all streamlines from `x` and `y`.

Value

- A `dist` object of size $x@n_streamlines$ when `y` is `NULL` or a `bundle`. The lower triangle stores all pairwise symmetric Hausdorff distances (computed in C++). Use `as.matrix()` to obtain the full $n \times n$ matrix.
- A numeric vector of length $x@n_streamlines$ when `y` is a `streamline`.

See Also

[compute_hausdorff_distance\(\)](#)

Examples

```
s11 <- streamline(points = cbind(X = runif(10), Y = runif(10), Z = runif(10)))
s12 <- streamline(points = cbind(X = runif(10), Y = runif(10), Z = runif(10)))
b <- bundle(streamlines = list(s11, s12))

# pairwise dist object (size 2)
compute_hausdorff_distance(b)
as.matrix(compute_hausdorff_distance(b))

# distances from s11 to each streamline in b
compute_hausdorff_distance(b, s11)
```

compute_hausdorff_distance-fiber-streamline-method

[compute_hausdorff_distance\(\)](#) *method for two streamline objects*

Description

[compute_hausdorff_distance\(\)](#) method for two streamline objects

Arguments

`x` A `streamline` object.
`y` A `streamline` object.

Value

A non-negative numeric scalar equal to $\max(d_H(x \rightarrow y), d_H(y \rightarrow x))$, where $d_H(A \rightarrow B) = \max_{a \in A} \min_{b \in B} \|a - b\|_2$ is the directed Hausdorff distance. The core computation is performed in C++ via `hausdorff_distance_cpp()`.

See Also

`compute_hausdorff_distance()`

Examples

```
s11 <- streamline(points = cbind(X = runif(10), Y = runif(10), Z = runif(10)))
s12 <- streamline(points = cbind(X = runif(10), Y = runif(10), Z = runif(10)))
compute_hausdorff_distance(s11, s12)
```

get_curvature

Curvature of a streamline

Description

`get_curvature()` is function that computes the curvature of a [streamline](#) object. The curvature $\kappa(s)$ at each point along the arc-length abscissa is computed using cubic smoothing splines (3 degrees of freedom per component).

Usage

```
get_curvature(x)
```

Arguments

`x` A [streamline](#) object.

Value

A non-negative numeric vector of length `x@n_points` giving the curvature $\kappa(s)$ at each sampled point along the streamline. Higher values indicate sharper bending at that location.

Examples

```
s1 <- streamline(points = cbind(X = runif(10), Y = runif(10), Z = runif(10)))
get_curvature(s1)
```

get_curvilinear_length *Curvilinear length of a streamline*

Description

get_curvilinear_length() is a function that computes the total arc-length of a [streamline](#) object as the sum of Euclidean segment lengths between consecutive points.

Usage

```
get_curvilinear_length(x)
```

Arguments

x A [streamline](#) object.

Value

A non-negative numeric scalar.

Examples

```
s1 <- streamline(points = cbind(X = runif(10), Y = runif(10), Z = runif(10)))
get_curvilinear_length(s1)
```

get_euclidean_length *Euclidean length of a streamline*

Description

get_euclidean_length() is a function that computes the Euclidean (straight-line) distance of a [streamline](#) object.

Usage

```
get_euclidean_length(x)
```

Arguments

x A [streamline](#) object.

Value

A non-negative numeric scalar.

Examples

```
sl <- streamline(points = cbind(X = runif(10), Y = runif(10), Z = runif(10)))  
get_euclidean_length(sl)
```

get_sinusosity	<i>Sinusosity of a streamline</i>
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Description

get_sinusosity() is a function that computes the ratio of curvilinear length to Euclidean length for a [streamline](#) object, with a value of 1 indicating a perfectly straight streamline and larger values indicating greater curviness.

Usage

```
get_sinusosity(x)
```

Arguments

x A [streamline](#) object.

Value

A numeric scalar ≥ 1 .

Examples

```
sl <- streamline(points = cbind(X = runif(10), Y = runif(10), Z = runif(10)))  
get_sinusosity(sl)
```

get_torsion	<i>Torsion of a streamline</i>
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Description

get_torsion() is function that computes the torsion of a [streamline](#) object. The torsion $\tau(s)$ at each point along the arc-length abscissa is computed using cubic smoothing splines (4 degrees of freedom per component).

Usage

```
get_torsion(x)
```

Arguments

x A [streamline](#) object.

Value

A numeric vector of length `x@n_points` giving the torsion $\tau(s)$ at each sampled point along the streamline. Positive values indicate right-handed twisting; negative values indicate left-handed twisting; zero indicates a planar curve at that location.

Examples

```
s1 <- streamline(points = cbind(X = runif(10), Y = runif(10), Z = runif(10)))
get_torsion(s1)
```

is_bundle	<i>Test whether an object is a bundle</i>
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Description

Test whether an object is a bundle

Usage

```
is_bundle(x)
```

Arguments

x An object.

Value

TRUE if x is of class `bundle`, otherwise FALSE.

Examples

```
s1 <- streamline(points = cbind(X = 1:5, Y = 1:5, Z = 1:5))
b <- bundle(streamlines = list(s1))
is_bundle(b)    # TRUE
is_bundle(s1)  # FALSE
```

is_bundle_set	<i>Test whether an object is a bundle_set</i>
---------------	---

Description

Test whether an object is a `bundle_set`

Usage

```
is_bundle_set(x)
```

Arguments

x An object.

Value

TRUE if x is of class `bundle_set`, otherwise FALSE.

Examples

```
s1 <- streamline(points = cbind(X = 1:5, Y = 1:5, Z = 1:5))
b <- bundle(streamlines = list(s1))
bs <- bundle_set(bundles = list(b))
is_bundle_set(bs) # TRUE
is_bundle_set(b)  # FALSE
```

is_streamline	<i>Test whether an object is a streamline</i>
---------------	---

Description

Test whether an object is a `streamline`

Usage

```
is_streamline(x)
```

Arguments

x An object.

Value

TRUE if x is of class `streamline`, otherwise FALSE.

Examples

```
s1 <- streamline(points = cbind(X = 1:5, Y = 1:5, Z = 1:5))
is_streamline(s1) # TRUE
is_streamline(42) # FALSE
```

reparametrize

Reparametrize a streamline or bundle onto a uniform arc-length grid

Description

reparametrize() is an S7 generic that resamples the 3-D coordinates (and any numeric @point_data attributes) of a tractography object onto a uniform arc-length grid using linear interpolation, with methods available for the following classes:

- `fiber::bundle`
- `fiber::streamline`

Usage

```
reparametrize(x, n_points = NULL)
```

Arguments

`x` A [streamline](#) or [bundle](#) object.

`n_points` Number of equally-spaced arc-length points to use.

- For a single [streamline](#), defaults to `x@n_points`.
- For a [bundle](#), defaults to the rounded mean number of points across all streamlines.

Pass NULL to use these defaults explicitly.

Value

An object of the same class as `x` reparametrized onto the new grid.

Examples

```
# reparametrize a single streamline to 10 points
s1 <- streamline(points = cbind(X = runif(5), Y = runif(5), Z = runif(5)))
s1_reparam <- reparametrize(s1, n_points = 10)
# reparametrize a bundle to the mean number of points across its streamlines
s11 <- streamline(points = cbind(X = runif(5), Y = runif(5), Z = runif(5)))
s12 <- streamline(points = cbind(X = runif(10), Y = runif(10), Z = runif(10)))
b <- bundle(streamlines = list(s11, s12))
bundle_reparam <- reparametrize(b)
```

```
reparametrize-fiber-bundle-method
      reparametrize() method for bundle objects
```

Description

Resamples every [streamline](#) inside a [bundle](#) onto a common uniform arc-length grid. See [reparametrize\(\)](#) for the full parameter documentation.

Arguments

<code>x</code>	A bundle object.
<code>n_points</code>	Number of equally-spaced arc-length points to use. <ul style="list-style-type: none"> • For a single streamline, defaults to <code>x@n_points</code>. • For a bundle, defaults to the rounded mean number of points across all streamlines. <p>Pass NULL to use these defaults explicitly.</p>

Value

A [bundle](#) reparametrized onto the new grid. Every streamline in the returned bundle has exactly `n_points` rows in `@points` (defaulting to the rounded mean number of points across all streamlines when `n_points` is NULL). `@streamline_data` and `@bundle_data` are preserved unchanged.

See Also

[reparametrize\(\)](#)

Examples

```
s11 <- streamline(points = cbind(X = runif(5), Y = runif(5), Z = runif(5)))
s12 <- streamline(points = cbind(X = runif(10), Y = runif(10), Z = runif(10)))
b <- bundle(streamlines = list(s11, s12))
b_reparam <- reparametrize(b, n_points = 8)
b_reparam[[1]]@n_points # 8
b_reparam[[2]]@n_points # 8
```

reparametrize-fiber-streamline-method
[reparametrize\(\)](#) *method for streamline objects*

Description

Resamples a single [streamline](#) onto a uniform arc-length grid. See [reparametrize\(\)](#) for the full parameter documentation.

Arguments

- | | |
|-----------------------|--|
| <code>x</code> | A streamline object. |
| <code>n_points</code> | Number of equally-spaced arc-length points to use. <ul style="list-style-type: none">• For a single streamline, defaults to <code>x@n_points</code>.• For a bundle, defaults to the rounded mean number of points across all streamlines. Pass NULL to use these defaults explicitly. |

Value

A [streamline](#) reparametrized onto the new grid. The returned object has `@points` resampled to exactly `n_points` rows via linear interpolation, and any numeric `@point_data` entries likewise resampled. Non-numeric `@point_data` entries are dropped with a warning. `@streamline_data` is preserved unchanged.

See Also

[reparametrize\(\)](#)

Examples

```
s1 <- streamline(  
  points = cbind(X = runif(10), Y = runif(10), Z = runif(10)),  
  point_data = list(FA = runif(10))  
)  
s1_reparam <- reparametrize(s1, n_points = 20)  
s1_reparam@n_points # 20
```

streamline

*Streamline S7 class***Description**

A streamline represents a single fibre tract. It stores three data compartments:

- `@points` — a $P \times 3$ numeric matrix with column names "X", "Y", and "Z" holding the 3-D coordinates of the P points along the tract.
- `@point_data` — a named list of numeric vectors, each of length P , holding additional per-point scalar attributes (e.g. fractional anisotropy). Coordinates are **not** stored here.
- `@streamline_data` — a named list of scalars (length-1 values of any type) holding per-streamline attributes (e.g. a tract-level weight or mean FA, or a character label).

Usage

```
streamline(points = NULL, point_data = list(), streamline_data = list())
```

Arguments

`points` A $P \times 3$ numeric matrix with column names "X", "Y", and "Z" giving the 3-D coordinates of the streamline points.

`point_data` A named list of numeric vectors, each of length P , holding additional per-point attributes. Must **not** include "X", "Y", or "Z" (those live in `@points`).

`streamline_data` A named list of per-streamline scalars (length-1, any type).

Value

A streamline S7 object.

Methods for standard generics

The following methods are defined for streamline objects:

- `format(x, ...)`: Returns a cli-formatted string describing the streamline object.
- `print(x, ...)`: Prints the formatted string to the console and invisibly returns `x`.

Additional properties

`@n_points` An integer scalar giving the number of points in the streamline (read-only).

`@point_attributes` A character vector of the names of the per-point attributes stored in `@point_data` (read-only).

`@streamline_attributes` A character vector of the names of the per-streamline attributes (read-only).

Examples

```
# Create a streamline with 5 points and some attributes
sl <- streamline(
  points = cbind(
    X = c(0, 1, 1, 1, 0),
    Y = c(0, 0, 1, 1, 1),
    Z = c(0, 0, 0, 1, 1)
  ),
  point_data = list(FA = c(0.5, 0.6, 0.7, 0.8, 0.9)),
  streamline_data = list(mean_FA = 0.7, label = "CST")
)
sl@n_points          # 5
sl@point_attributes # "FA"
sl@streamline_attributes # c("mean_FA", "label")

# format() and print() methods
format(sl)
print(sl)
```


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