Package: pavdata (via r-universe)

June 20, 2026

Title Transportation Infrastructure Data Toolbox

Version 0.1.0

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Description An open-source toolbox for storing, validating, managing, and exploring transportation infrastructure data. Provides a relational data model for binders, aggregates, mixtures, and test results, with a human-readable file format (.pavdata) aligned with FAIR principles.

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Encoding UTF-8

Depends R (>= 4.0)

Imports jsonlite, uuid, graphics

Suggests testthat (>= 3.0.0)

Config/testthat/edition 3

Config/roxygen2/version 8.0.0

NeedsCompilation no

Repository https://vilmarfausttino.r-universe.dev

Date/Publication 2026-06-19 18:11:51 UTC

RemoteUrl https://github.com/cran/pavdata

RemoteRef HEAD

RemoteSha 5efc55153efa445a6fe3c57a25e2ae3c1d2292bf

Contents

`is.pavdata` *Test whether an object is a PavData object*

Description

Test whether an object is a PavData object

Usage

```
is.pavdata(x)
```

Arguments

`x`	Any R object.

Value

A single logical value: `TRUE` if `x` is a PavData object, `FALSE` otherwise.

Examples

```
is.pavdata(pav_new("sample", name = "Am_001"))
```

pav_check *Validate a single PavData object*

Description

Checks a PavData object for required fields (level 1) and for values outside physically plausible ranges (level 2). Missing optional fields do not cause failure.

Usage

```
pav_check(x, verbose = TRUE)
```

Arguments

x	A PavData object created by pav_new.
verbose	Logical; if TRUE (default), prints the validation outcome and any issues.

Value

Invisibly, a list with valid (logical) and issues (a character vector of messages).

Examples

```
pav_check(pav_new("binder", name = "CAP 50/70", binder_type = "CAP 50/70"))
```

pav_check_integrity *Check relational integrity of a collection*

Description

Validates every object in a collection and verifies that reference fields (foreign keys) point to existing objects, reporting failed objects and broken references.

Usage

```
pav_check_integrity(objects, verbose = TRUE)
```

Arguments

objects	A named list of PavData objects, as returned by pav_read.
verbose	Logical; if TRUE (default), prints a summary of the check.

Value

Invisibly, a list with valid (logical), total, failed and broken_refs.

Examples

```
b <- pav_new("binder", name = "CAP 50/70", binder_type = "CAP 50/70")
pav_check_integrity(list(b))
```

`pav_library` *Create a PavData library*

Description

Creates an empty in-memory library to store, index and query PavData objects.

Usage

```
pav_library()
```

Value

A `pav_library` object (an environment) with an empty object store and type index.

Examples

```
lib <- pav_library()
```

`pav_library_load` *Load a .pavdata file into a library*

Description

Reads a `.pavdata` file and adds all of its objects to an existing library, updating the type index.

Usage

```
pav_library_load(lib, path)
```

Arguments

`lib`	A `pav_library` object created by `pav_library`.
`path`	Character string with the path to a `.pavdata` file.

Value

The library `lib`, invisibly.

Examples

```
b <- pav_new("binder", name = "CAP 50/70", binder_type = "CAP 50/70")
tmp <- tempfile(fileext = ".pavdata")
pav_write(b, tmp)
lib <- pav_library()
pav_library_load(lib, tmp)
```

`pav_list` *List objects in a library*

Description

Lists objects stored in a library, optionally filtered by type, by a substring of the name, or by binder type, up to a maximum number of results.

Usage

```
pav_list(
  lib,
  type = NULL,
  name_contains = NULL,
  binder_type = NULL,
  limit = 20
)
```

Arguments

`lib`	A `pav_library` object created by `pav_library`.
`type`	Character string with an object type to filter by, or `NULL` (default) for all types.
`name_contains`	Character string matched against object names (case-insensitive), or `NULL` (default).
`binder_type`	Character string with a binder type to filter by, or `NULL` (default).
`limit`	Integer; maximum number of objects to return (default 20).

Value

Invisibly, a list with the matching objects.

Examples

```
b <- pav_new("binder", name = "CAP 50/70", binder_type = "CAP 50/70")
tmp <- tempfile(fileext = ".pavdata")
pav_write(b, tmp)
lib <- pav_library()
pav_library_load(lib, tmp)
pav_list(lib, type = "binder")
```

pav_load *Read and validate a .pavdata file*

Description

Reads a `.pavdata` file with `pav_read` and then validates each object, warning if any fail validation.

Usage

```
pav_load(path)
```

Arguments

`path`	Character string with the path to a `.pavdata` file.

Value

A named list of PavData objects, keyed by object id.

Examples

```
b <- pav_new("binder", name = "CAP 50/70", binder_type = "CAP 50/70")
tmp <- tempfile(fileext = ".pavdata")
pav_write(b, tmp)
objects <- pav_load(tmp)
```

pav_new *Create a PavData object*

Description

Constructs a PavData object of a given type, attaching the common metadata (id, name, type, version, creation timestamp, source, references, notes and history) shared by all object types.

Usage

```
pav_new(object_type, ...)
```

Arguments

`object_type`	Character string with the object type. One of `"sample"`, `"binder"`, `"aggregate"`, `"mixture"`, `"binder_test"`, `"aggregate_test"`, `"mixture_test"` or `"reference"`.
`...`	Named arguments with the type-specific fields (for example, `binder_type` for a binder, or `binder_id` and `aggregate_id` for a mixture).

Value

A PavData object: a list with S3 classes `pavdata_<type>`, `pavdata_object` and `list`.

Examples

```
b <- pav_new("binder", name = "CAP 50/70", binder_type = "CAP 50/70")
m <- pav_new("mixture", name = "Mixture 1", binder_id = b$id,
             aggregate_id = "agg_001")
```

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pav_read *Read a .pavdata file*

Description

Reads a `.pavdata` file and reconstructs the PavData objects, restoring their S3 classes and indexing them by id.

Usage

```
pav_read(path)
```

Arguments

`path`	Character string with the path to a `.pavdata` file.

Value

A named list of PavData objects, keyed by object id.

Examples

```
b <- pav_new("binder", name = "CAP 50/70", binder_type = "CAP 50/70")
tmp <- tempfile(fileext = ".pavdata")
pav_write(b, tmp)
objects <- pav_read(tmp)
```

pav_view *View the details of an object*

Description

Prints the metadata and the filled fields of a single object stored in a library, including its non-empty nested values.

Usage

```
pav_view(lib, id)
```

Arguments

lib	A pav_library object created by pav_library.
id	Character string with the id of the object to view.

Value

Invisibly, the object, or NULL if it is not found.

Examples

```
b <- pav_new("binder", name = "CAP 50/70", binder_type = "CAP 50/70")
tmp <- tempfile(fileext = ".pavdata")
pav_write(b, tmp)
lib <- pav_library()
pav_library_load(lib, tmp)
pav_view(lib, b$id)
```

pav_write *Write PavData objects to a .pavdata file*

Description

Serializes one or more PavData objects to a .pavdata file (JSON content), grouping them by type. The .pavdata extension is appended when missing.

Usage

```
pav_write(x, path, pretty = TRUE)
```

Arguments

x	A single PavData object or a list of PavData objects.
path	Character string with the output file path.
pretty	Logical; if TRUE (default), the JSON is indented for readability.

Value

The output path, invisibly.

Examples

```
b <- pav_new("binder", name = "CAP 50/70", binder_type = "CAP 50/70")
tmp <- tempfile(fileext = ".pavdata")
pav_write(b, tmp)
```

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pavdata_sample_path *Path to the bundled example dataset*

Description

Returns the file path to the example .pavdata dataset shipped with the package (296 Brazilian asphalt mixture records).

Usage

```
pavdata_sample_path()
```

Value

A character string with the full path to the dataset file, or an empty string if the package is not installed.

Examples

```
path <- pavdata_sample_path()
```

pavdata_type *Get the type of a PavData object*

Description

Get the type of a PavData object

Usage

```
pavdata_type(x)
```

Arguments

x	A PavData object created by pav_new.

Value

A character string with the object type (for example, "binder"), taken from its S3 class.

Examples

```
pavdata_type(pav_new("binder", name = "CAP 50/70", binder_type = "CAP 50/70"))
```

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