

# Package ‘neonstore’

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**Title** NEON Data Store

**Version** 0.5.1

**Description** The National Ecological Observatory Network (NEON) provides access to its numerous data products through its REST API, [<https://data.neonscience.org/data-api/>](https://data.neonscience.org/data-api/). This package provides a high-level user interface for downloading and storing NEON data products. Unlike 'neonUtilities', this package will avoid repeated downloading, provides persistent storage, and improves performance. 'neonstore' can also construct a local 'duckdb' database of stacked tables, making it possible to work with tables that are far to big to fit into memory.

**License** MIT + file LICENSE

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neon_citation	<i>Generate the appropriate citation for your data</i>
---------------	--

---

**Description**

Generate the appropriate citation for your data

**Usage**

```
neon_citation(product = NULL, download_date = Sys.Date(), dir = neon_dir())
```

## Arguments

product	A NEON productCode or list of product codes, see examples.
download_date	Date of download to be included in citation. default is today's date, see details.
dir	Location where files should be downloaded. By default will use the appropriate applications directory for your system (see <code>tools::R_user_dir()</code> ). This default also be configured by setting the environmental variable NEONSTORE_HOME, see <a href="#">Sys.setenv</a> or <a href="#">Renviron</a> .

## Details

Note that the `neon_download()` does not record download date for each file. Citing a single product download date is after all rather meaningless, as parts of a products may not have all been downloaded on different dates. Indeed, `neon_download()` is designed in precisely this way, to allow easy updating of downloads without re-downloading older data.

## Value

returns a `utils::bibentry` object, which can be used as text or formatted for bibtex.

## References

<https://www.neonscience.org/data-samples/data-policies-citation>

## Examples

```
# may be slow
neon_citation("DP1.10003.001")

## or the citation for all products in store:
neon_citation()

## as bibtex
format(neon_citation("DP1.10003.001"), "bibtex")
```

---

neon\_cloud

*neon cloud*

---

## Description

neon cloud

**Usage**

```

neon_cloud(
  table,
  product,
  start_date = NA,
  end_date = NA,
  site = NA,
  type = "basic",
  release = NA,
  quiet = FALSE,
  api = "https://data.neonscience.org/api/v0",
  unify_schemas = FALSE,
  .token = Sys.getenv("NEON_TOKEN")
)

```

**Arguments**

table	NEON table name
product	A NEON productCode or list of product codes, see examples.
start_date	Download only files as recent as (YYYY-MM-DD). Leave as NA to download up to the most recent available data.
end_date	Download only files up to end_date (YYYY-MM-DD). Leave as NA to download all prior data.
site	4-letter site code(s) to filter on. Leave as NA to search all.
type	Should we prefer the basic or expanded version of this product? Note that not all products have expanded formats.
release	Select only data files associated with a particular release tag, see <a href="https://www.neonscience.org/data-samples/data-management/data-revisions-releases">https://www.neonscience.org/data-samples/data-management/data-revisions-releases</a> , e.g. "RELEASE-2021". Releases are associated with a specific DOI and the promise that files associated with a particular release will not change.
quiet	Should download progress be displayed?
api	the URL to the NEON API, leave as default.
unify_schemas	if cloud-read fails to collect data due to miss-matched schemas, set this to TRUE. Warning: Results in much slower reads and may demand more memory due to parsing the schema of each file, especially when many files are involved.
.token	an authentication token from NEON. A token is not required but will allow access to a higher number of requests before rate limiting applies, see <a href="https://data.neonscience.org/data-api/rate-limiting/#api-tokens">https://data.neonscience.org/data-api/rate-limiting/#api-tokens</a> . Note that once files are downloaded once, neonstore provides persistent access to them without further interaction required with the API.

**Value**

lazy data frame

---

neon_data	<i>Query the NEON API for URLs of matching data products Repeated requests will be cached</i>
-----------	---

---

### Description

Query the NEON API for URLs of matching data products Repeated requests will be cached

### Usage

```
neon_data(
  product,
  start_date = NA,
  end_date = NA,
  site = NA,
  type = NA,
  release = NA,
  quiet = FALSE,
  api = "https://data.neonscience.org/api/v0",
  .token = Sys.getenv("NEON_TOKEN")
)
```

### Arguments

product	A NEON productCode or list of product codes, see examples.
start_date	Download only files as recent as (YYYY-MM-DD). Leave as NA to download up to the most recent available data.
end_date	Download only files up to end_date (YYYY-MM-DD). Leave as NA to download all prior data.
site	4-letter site code(s) to filter on. Leave as NA to search all.
type	Should we prefer the basic or expanded version of this product? Note that not all products have expanded formats.
release	Select only data files associated with a particular release tag, see <a href="https://www.neonscience.org/data-samples/data-management/data-revisions-releases">https://www.neonscience.org/data-samples/data-management/data-revisions-releases</a> , e.g. "RELEASE-2021". Releases are associated with a specific DOI and the promise that files associated with a particular release will not change.
quiet	Should download progress be displayed?
api	the URL to the NEON API, leave as default.
.token	an authentication token from NEON. A token is not required but will allow access to a higher number of requests before rate limiting applies, see <a href="https://data.neonscience.org/data-api/rate-limiting/#api-tokens">https://data.neonscience.org/data-api/rate-limiting/#api-tokens</a> . Note that once files are downloaded once, neonstore provides persistent access to them without further interaction required with the API.

**Value**

a data.frame containing the name, filesize (in bytes), checksums (columns md5, crc32, or crc32c, though each product will use only one of these), url, and release status.

**Examples**

```
x <- neon_data("DP1.10003.001")
x <- neon_data("DP1.10003.001", release="RELEASE-2021")
```

---

neon\_db

*Cache-able duckdb database connection*


---

**Description**

Cache-able duckdb database connection

**Usage**

```
neon_db(
  dir = neon_db_dir(),
  read_only = TRUE,
  memory_limit = getOption("duckdb_memory_limit", NA),
  ...
)
```

**Arguments**

<code>dir</code>	Location where files should be downloaded. By default will use the appropriate applications directory for your system (see <code>tools::R_user_dir()</code> ). This default also be configured by setting the environmental variable <code>NEONSTORE_HOME</code> , see <a href="#">Sys.setenv</a> or <a href="#">Renviron</a> .
<code>read_only</code>	allow concurrent connections by enforcing <code>read_only</code> . See details.
<code>memory_limit</code>	Set a memory limit for duckdb, in GB. This can also be set for the session by using options, e.g. <code>options(duckdb_memory_limit=10)</code> for a limit of 10GB. On most systems duckdb will automatically set a limit to 80% of machine capacity if not set explicitly.
<code>...</code>	additional arguments to <code>dbConnect</code>

**Details**

Creates a connection to a permanent duckdb database instance in the provided directory (see [neon\\_dir\(\)](#)). This connection is also cached, so that code which repeatedly calls `[neon_db]` will not stall or hang. Only `read_only` connections will be cached.

NOTE: `[duckdb:::duckdb()]` can only support a single read-write connection at a time. The default option of `read_only = TRUE` allows multiple connections. `[neon_store()]` will automatically set this to `FALSE` to allow data import.

**Examples**

```
# tempfile used for illustration only
neon_db(tempfile())
```

---

neon_db_dir	<i>Default directory for persistent NEON database</i>
-------------	---

---

**Description**

Use `neon_db_dir()` to view or access the currently active database directory. By default, this uses the appropriate application directory for your operating system, see [tools::R\\_user\\_dir\(\)](#). This location can be overridden by setting the environmental variable `NEONSTORE_DB`.

**Usage**

```
neon_db_dir()
```

**Value**

the active neonstore directory.

**Examples**

```
neon_db_dir()

## Override with an environmental variable:
Sys.setenv(NEONSTORE_DB = tempdir())
neon_db_dir()
## Unset
Sys.unsetenv("NEONSTORE_DB")
```

---

neon_delete_db	<i>delete the local NEON database</i>
----------------	---------------------------------------

---

**Description**

delete the local NEON database

**Usage**

```
neon_delete_db(db_dir = neon_db_dir(), ask = interactive())
```

**Arguments**

db_dir	neon database location (configurable with the NEONSTORE_DB environmental variable)
ask	Ask for confirmation first?

**Details**

Just a helper function that deletes the NEON database files, which are found under `file.path(neon_dir(), "database")`. This does not delete downloaded raw data, which can easily be re-loaded with `neon_store()`. Usually unnecessary but can be helpful in resetting a corrupt database.

If you want to delete all raw data files downloaded by neonstore as well, simply delete the entire directory given by `neon_dir()`

**Examples**

```
# Create a db
dir <- tempfile()
db <- neon_db(dir)

# Delete it
neon_delete_db(dir, ask = FALSE)
```

---

neon_dir	<i>Default directory for persistent NEON file store</i>
----------	---

---

**Description**

Use `neon_dir()` to view or access the currently active local store. By default, `neon_download()` downloads files into the `neon_dir()`, which uses an appropriate application directory for your operating system, see `tools::R_user_dir()`. This location can be overridden by setting the environmental variable `NEONSTORE_HOME`. neonstore functions (e.g. `neon_index()`, and `neon_read()`) look for files in the `neon_dir()` directory by default. (All functions can also take a one-off argument to `dir` in the function call in place of the calling `neon_dir()` to access the default.



**Usage**

```
neon_dir()
```

**Value**

the active neonstore directory.

**Examples**

```
neon_dir()

## Override with an environmental variable:
Sys.setenv(NEONSTORE_HOME = tempdir())
neon_dir()
## Unset
Sys.unsetenv("NEONSTORE_HOME")
```

---

neon_disconnect	<i>Disconnect from the neon database</i>
-----------------	--

---

**Description**

Disconnect from the neon database

**Usage**

```
neon_disconnect(db = neon_db())
```

**Arguments**

db                    link to an existing database connection

---

neon_download	<i>Download NEON data products into a local store</i>
---------------	---

---

**Description**

Download NEON data products into a local store

**Usage**

```

neon_download(
  product,
  table = NA,
  site = NA,
  start_date = NA,
  end_date = NA,
  type = "basic",
  release = NA,
  quiet = FALSE,
  verify = TRUE,
  unique = TRUE,
  dir = neon_dir(),
  get_zip = FALSE,
  unzip = FALSE,
  api = "https://data.neonscience.org/api/v0",
  .token = Sys.getenv("NEON_TOKEN")
)

```

**Arguments**

product	A NEON productCode or list of product codes, see examples.
table	Include only files matching this table name (or regex pattern). (optional).
site	4-letter site code(s) to filter on. Leave as NA to search all.
start_date	Download only files as recent as (YYYY-MM-DD). Leave as NA to download up to the most recent available data.
end_date	Download only files up to end_date (YYYY-MM-DD). Leave as NA to download all prior data.
type	Should we prefer the basic or expanded version of this product? Note that not all products have expanded formats.
release	Select only data files associated with a particular release tag, see <a href="https://www.neonscience.org/data-samples/data-management/data-revisions-releases">https://www.neonscience.org/data-samples/data-management/data-revisions-releases</a> , e.g. "RELEASE-2021". Releases are associated with a specific DOI and the promise that files associated with a particular release will not change.
quiet	Should download progress be displayed?
verify	Should downloaded files be compared against the MD5 hash reported by the NEON API to verify integrity? (default TRUE)
unique	Should we skip downloads of files we already have? Note: file comparisons are based on file hash, which will omit files that have identical content but different names.
dir	Location where files should be downloaded. By default will use the appropriate applications directory for your system (see <code>tools::R_user_dir()</code> ). This default also be configured by setting the environmental variable NEONSTORE_HOME, see <a href="#">Sys.setenv</a> or <a href="#">Renviron</a> .

get_zip	should we attempt to download .zip archive versions of files? default FALSE, as zip archives are being deprecated from NEON API starting in early 2021.
unzip	should we extract .zip files? (default TRUE). Note: .zip files are preserved in the store to avoid repeated downloads. Use of .zip files in NEON API is now deprecated in favor of requesting individual files.
api	the URL to the NEON API, leave as default.
.token	an authentication token from NEON. A token is not required but will allow access to a higher number of requests before rate limiting applies, see <a href="https://data.neonscience.org/data-api/rate-limiting/#api-tokens">https://data.neonscience.org/data-api/rate-limiting/#api-tokens</a> . Note that once files are downloaded once, neonstore provides persistent access to them without further interaction required with the API.

## Details

Each NEON data product consists of a collection of objects (e.g. tables), which are in turn broken into individual files by site and sampling month. Additionally, many NEON products have been expanded, including some additional columns. Consequently, users must specify if they want the "basic" or "expanded" version of this data.

In the products table (see [neon\\_products](#)), the productHasExpanded column indicates if the data product has expanded, and the columns productHasBasicDescription and productHasExpandedDescription provide a detailed explanation of the differences between the "expanded" and "basic" versions of that particular product.

The API allows users to request component files directly. By default, `neon_download()` will download all available extensions. Users can request only products of a certain format (e.g. .csv or .h5) by altering the `file_regex` argument (see examples).

Prior to 2021, the API provided access to a .zip file containing all the component objects (e.g. tables) for that product at that site and sampling month.

`neon_download()` will avoid downloading metadata files which are bitwise identical to other files in the same download request, as indicated by the `crc32` hash reported by the API. These typically include metadata that are shared across the product as a whole, but are for some reason included in each sampling month for each site – potentially thousands of duplicates. These duplicates are also packaged within the .zip downloads where it is not possible to exclude them from the download.

## Examples

```
## Omit dir=tempfile() to use persistent storage
neon_download("DP1.10003.001",
              start_date = "2018-01-01",
              end_date = "2019-01-01",
              site = "YELL",
              dir = tempfile())

## Advanced use: filter for a particular table in the product
neon_download(product = "DP1.10003.001",
              start_date = "2018-01-01",
              end_date = "2019-01-01",
```

```

site = "YELL",
table = "countdata",
dir = tempfile()

```

---

neon\_export

*export local neon store as a zip archive*


---

### Description

Export all or select files from your neon store as a zip archive. This can be useful if you want to bypass accessing the API, such as for archiving the files required for your analysis so that they can be re-created by other users without an API key, or without waiting for the individual download, or any other time you want to share or distribute your local store.

### Usage

```

neon_export(
  archive = paste(Sys.Date(), "neonstore.zip", sep = "-"),
  product = NA,
  table = NA,
  site = NA,
  start_date = NA,
  end_date = NA,
  type = NA,
  ext = NA,
  timestamp = NA,
  hash = NULL,
  dir = neon_dir()
)

```

### Arguments

archive	path to the zip archive to be created.#'
product	A NEON productCode or list of product codes, see examples.
table	Include only files matching this table name (or regex pattern). (optional).
site	4-letter site code(s) to filter on. Leave as NA to search all.
start_date	Download only files as recent as (YYYY-MM-DD). Leave as NA to download up to the most recent available data.
end_date	Download only files up to end_date (YYYY-MM-DD). Leave as NA to download all prior data.
type	Should we prefer the basic or expanded version of this product? Note that not all products have expanded formats.
ext	only match files with this file extension(s)

timestamp	only match timestamps prior this. See details in <a href="#">neon_index()</a> . Should be a datetime POSIXct object (or coerce-able string)
hash	name of a hashing algorithm to check file integrity. Can be "md5", "sha1", or "sha256" currently; or set to <b>NULL</b> (default) to skip hash computation.
dir	Location where files should be downloaded. By default will use the appropriate applications directory for your system (see <a href="#">tools::R_user_dir()</a> ). This default also be configured by setting the environmental variable NEONSTORE_HOME, see <a href="#">Sys.setenv</a> or <a href="#">Renviron</a> .

**Value**

table of selected files and metadata, from [neon\\_index\(\)](#), invisibly.

**See Also**

[neon\\_import\(\)](#), [neon\\_citation\(\)](#)

**Examples**

```
archive <- tempfile()
dir <- tempdir()
neon_export(archive, dir = dir)
```

---

neon_export_db	<i>Export NEON database to parquet</i>
----------------	--

---

**Description**

Export your current database. This can be important to (1) archive and share your database files with another user or machine, (2) expose your database using an S3 bucket using [neon\\_remote\\_db\(\)](#), (3) assist in upgrading your duckdb version.

**Usage**

```
neon_export_db(dir = file.path(neon_dir(), "parquet"), db = neon_db())
```

**Arguments**

dir	directory to which parquet export is written.
db	Connection to your local NEON database

---

neon\_filename\_parser *NEON filename parser*

---

### Description

Parse filenames into their component metadata. See details for definition of each metadata field, or consult the NEON documentation linked below. <https://data.neonscience.org/file-naming-conventions>

### Usage

```
neon_filename_parser(x)
```

### Arguments

x                      vector of NEON filenames

### Details

#### Metadata components::

- NEON A four-character alphanumeric code, denoting the organizational origin of the data product and identifying the product as operational; data collected as part of a special data collection exercise are designated by a separate, unique alphanumeric code created by the PI.
- DOM A three-character alphanumeric code, referring to the domain of data acquisition (D01 - D20).
- SITE A four-character alphanumeric code, referring to the site of data acquisition; all sites are designated by a standardized four-character alphabetic code.
- DPL A three-character alphanumeric code, referring to data product processing level.
- PRNUM A five-character numeric code, referring to the data product number (see the Data Product Catalog at <http://data.neonscience.org/data-product-catalog>).
- REV A three-digit designation, referring to the revision number of the data product. The REV value is incremented by 1 each time a major change is made in instrumentation, data collection protocol, or data processing such that data from the preceding revision is not directly comparable to the new.
- HOR A three-character alphanumeric code for Spatial Index #1. Refers to measurement locations within one horizontal plane. For example, if five surface measurements were taken, one at each of the five soil array plots, the number in the HOR field would range from 001-005.
- VER A three-character alphanumeric code for Spatial Index #2. Refers to measurement locations within one vertical plane. For example, if eight temperature measurements are collected, one at each tower vertical level, the number in the VER field would range from 010-080.
- TMI A three-character alphanumeric code for the Temporal Index. Refers to the temporal representation, averaging period, or coverage of the data product (e.g., minute, hour, month, year, sub-hourly, day, lunar month, single instance, seasonal, annual, multi-annual). 000 = native resolution, 001 = native resolution or 1 minute, 002 = 2 minute, 005 = 5 minute, 015 = 15 minute, 030 = 30 minute, 060 = 60 minutes or 1 hour, 101-103 = native resolution

of replicate sensor 1, 2, and 3 respectively, 999 = Sensor conducts measurements at varied interval depending on air mass.

- DESC An abbreviated description of the data file or table.
- YYYY-MM Represents the year and month of the data in the file.
- PKGTYPE The type of data package downloaded. Options are 'basic', representing the basic download package, or 'expanded', representing the expanded download package (see more information below).
- GENTIME The date-time stamp when the file was generated, in UTC. The format of the date-time stamp is YYYYMMDDTHHmmSSZ.

#### **AOP Products Only (Airborne Observation Platform)::**

- FLHTDATE Date of flight, YYYYMMDD
- FLIGHTSTRT Start time of flight, YYYYMMDDHH
- FLHTSTRT Start time of flight, YMMDDHH
- IMAGEDATETIME Date and time of image capture, YYYYMMDDHHmmSS
- CCCCCC Digital camera serial number
- NNNN Sequential number for indexing files
- NNN Planned flightline number
- R Repeat number
- FFFFFFF Numeric code for an individual flightline
- EEEEEEE UTM easting of lower left corner
- NNNNNNN UTM northing of lower left corner

#### **Value**

a data frame in which filenames have been split into metadata components. Column names indicate the metadata field code, see details section for complete descriptions.

#### **References**

<https://data.neonscience.org/file-naming-conventions>

---

neon\_import

*Import a previously exported zip archive of raw NEON files*

---

#### **Description**

`neon_import()` only reads in previously saved archives from `neon_export()`. This can be useful in cases where see `neon_download()` to download data directly from NEON.

#### **Usage**

```
neon_import(archive, overwrite = TRUE, dir = neon_dir())
```

**Arguments**

archive	path to the zip archive to be imported
overwrite	should we overwrite any existing files?
dir	Location where files should be downloaded. By default will use the appropriate applications directory for your system (see <code>tools::R_user_dir()</code> ). This default also be configured by setting the environmental variable <code>NEONSTORE_HOME</code> , see <a href="#">Sys.setenv</a> or <a href="#">Renviron</a> .

**See Also**

[neon\\_export\(\)](#)

**Examples**

```
## tempfiles for example purposes only!
archive <- tempfile()
neon_dir <- tempdir()

neon_export(archive, dir = neon_dir)
neon_import(archive)
```

---

neon_import_db	<i>Import a NEON database exported from neon_export_db()</i>
----------------	--

---

**Description**

Import a NEON database exported from `neon_export_db()`

**Usage**

```
neon_import_db(
  dir = file.path(neon_dir(), "parquet"),
  db = neon_db(read_only = FALSE)
)
```

**Arguments**

dir	directory to which parquet export is written.
db	Connection to your local NEON database



---

 neon\_index

*Show information about all files downloaded to the local store*


---

## Description

NEON products consist of several individual components, which are in turn broken up by site and sampling month. By storing these individual files, neonstore enables more reproducible workflows that can be traced back to original, unaltered input data. These atomized files can be quickly and easily combined into unified tables, see [neon\\_read](#).

## Usage

```
neon_index(
  product = NA,
  table = NA,
  site = NA,
  start_date = NA,
  end_date = NA,
  type = NA,
  ext = NA,
  timestamp = NA,
  release = NA,
  hash = NULL,
  dir = neon_dir(),
  deprecated = TRUE
)
```

## Arguments

product	A NEON productCode or list of product codes, see examples.
table	Include only files matching this table name (or regex pattern). (optional).
site	4-letter site code(s) to filter on. Leave as NA to search all.
start_date	Download only files as recent as (YYYY-MM-DD). Leave as NA to download up to the most recent available data.
end_date	Download only files up to end_date (YYYY-MM-DD). Leave as NA to download all prior data.
type	Should we prefer the basic or expanded version of this product? Note that not all products have expanded formats.
ext	only match files with this file extension(s)
timestamp	only match timestamps prior this. See details in <a href="#">neon_index()</a> . Should be a datetime POSIXct object (or coerce-able string)
release	Select only data files associated with a particular release tag, see <a href="https://www.neonscience.org/data-samples/data-management/data-revisions-releases">https://www.neonscience.org/data-samples/data-management/data-revisions-releases</a> , e.g. "RELEASE-2021". Releases are associated with a specific DOI and the promise that files associated with a particular release will not change.

hash	name of a hashing algorithm to check file integrity. Can be "md5", "sha1", or "sha256" currently; or set to <code>NULL</code> (default) to skip hash computation.
dir	Location where files should be downloaded. By default will use the appropriate applications directory for your system (see <code>tools::R_user_dir()</code> ). This default also be configured by setting the environmental variable <code>NEONSTORE_HOME</code> , see <code>Sys.setenv</code> or <code>Renviron</code> .
deprecated	Should the index include files that have since been deprecated by more recent downloads? logical, default <code>TRUE</code> .

### Details

File names include metadata such as the file productCode, table name, site, and sampling month, as well as timestamp of creation. `neon_index()` parses this metadata from the file name string and returns the information in a convenient table, along with a path to each file.

**Regarding timestamps:** NEON will occasionally publish new versions of previously-released raw data files (which may or may not actually differ). The NEON download API, and hence `neon_download()`, only serve the most recent of such files, but earlier versions may still exist in your local neonstore if you downloaded them before the updated files were released. By default, `neon_read()` will always select the most recent of such files, thus avoiding duplication and providing the most updated data. For reproducibility however, it may be necessary to access older version instead. Setting the timestamp argument allows the user to filter out newer files and select the original ones instead. Unfortunately, at this time users cannot request the outdated data files from NEON API. For strict reproducibility, users should also archive their local store.

### See Also

`neon_download()`

### Examples

```
neon_index()

## Just bird survey product
neon_index("DP1.10003.001")
```

---

neon\_pane

*Open NEON database connection pane in RStudio*

---

### Description

This function launches the RStudio "Connection" pane to interactively explore the database.

**Usage**

```
neon_pane()
```

**Examples**

```
if (!is.null(getOption("connectionObserver"))) neon_pane()
```

---

neon_products	<i>Table of all NEON Data Products</i>
---------------	--

---

**Description**

Return a table of all NEON Data Products, including product descriptions and the productCode needed for [neon\\_download](#). (including list-columns).

**Usage**

```
neon_products(
  api = "https://data.neonscience.org/api/v0",
  .token = Sys.getenv("NEON_TOKEN")
)
```

**Arguments**

api	the URL to the NEON API, leave as default.
.token	an authentication token from NEON. A token is not required but will allow access to a higher number of requests before rate limiting applies, see <a href="https://data.neonscience.org/data-api/rate-limiting/#api-tokens">https://data.neonscience.org/data-api/rate-limiting/#api-tokens</a> . Note that once files are downloaded once, neonstore provides persistent access to them without further interaction required with the API.

**See Also**

[neon\\_download](#)

**Examples**

```
products <- neon_products()

# Or search for a keyword
i <- grepl("bird", products$keywords)
products[i, c("productCode", "productName")]
```

---

neon_read	<i>read in neon tabular data</i>
-----------	----------------------------------

---

### Description

read in neon tabular data

### Usage

```
neon_read(
  table = NA,
  product = NA,
  site = NA,
  start_date = NA,
  end_date = NA,
  ext = NA,
  timestamp = NA,
  release = NA,
  dir = neon_dir(),
  files = NULL,
  sensor_metadata = TRUE,
  keep_filename = FALSE,
  altrep = FALSE,
  ...
)
```

### Arguments

table	the name of a downloaded NEON table in the store, see <a href="#">neon_index</a>
product	A NEON productCode or list of product codes, see examples.
site	4-letter site code(s) to filter on. Leave as NA to search all.
start_date	Download only files as recent as (YYYY-MM-DD). Leave as NA to download up to the most recent available data.
end_date	Download only files up to end_date (YYYY-MM-DD). Leave as NA to download all prior data.
ext	only match files with this file extension(s)
timestamp	only match timestamps prior this. See details in <a href="#">neon_index()</a> . Should be a datetime POSIXct object (or coerce-able string)
release	Select only data files associated with a particular release tag, see <a href="https://www.neonscience.org/data-samples/data-management/data-revisions-releases">https://www.neonscience.org/data-samples/data-management/data-revisions-releases</a> , e.g. "RELEASE-2021". Releases are associated with a specific DOI and the promise that files associated with a particular release will not change.
dir	Location where files should be downloaded. By default will use the appropriate applications directory for your system (see <a href="#">tools::R_user_dir()</a> ). This default also be configured by setting the environmental variable NEONSTORE_HOME, see <a href="#">Sys.setenv</a> or <a href="#">Renviron</a> .

files	optionally, specify a vector of file paths directly (e.g. as provided from <code>neon_index</code> ) and specify table argument as <code>NULL</code> .
sensor_metadata	logical, default <code>TRUE</code> . Should we add metadata fields from file names of sensor data into the table? Adds <code>DomainID</code> , <code>SiteID</code> , <code>horizontalPosition</code> , <code>verticalPosition</code> , and <code>publicationDate</code> . Results in slower parsing.
keep_filename	Should we include a column indicating the original file name for each row? Can be a useful source of additional metadata that NEON may omit from the raw files (i.e. <code>siteID</code> ), but will also result in slower parsing. Default <code>FALSE</code> .
altrep	enable or disable <code>altrep</code> . Logical, default <code>FALSE</code> . Setting to <code>TRUE</code> can speed up reading, but may cause <code>vroom::vroom</code> to throw mapping error: Too many open files.
...	additional arguments to <code>vroom::vroom</code> , can usually be omitted.

## Details

NEON's tabular data files are separated out into separate `.csv` files for each site for each month of sampling. In principle, each file has identical columns. `vroom::vroom` can read in a data table that has been sharded into many files like this much much faster than other parsers can read in each table iteratively, (and thus can greatly out-perform the 'stacking' methods in `neonUtilities`).

When reading in very large numbers of files, it may be helpful to set `altrep = FALSE` to opt out of `vroom`'s fast `altrep` mechanism, which can cause `neon_read()` to fail when stacking thousands of files.

Unfortunately, not all datasets are entirely consistent in their use of columns. `neon_read` works around this by parsing such tables in groups of matching schema, which is still reasonably fast.

NEON sensor data products currently do not include important metadata columns containing `DomainID`, `SiteID`, `horizontalPosition`, `verticalPosition`, and `publicationDate` in the data files themselves, but only encode this in the in the raw file names. All though these values are shared across a raw data file, this information is lost when stacking the tables unless explicit columns are added to the data. This requires us to parse the files one-by-one, which is much slower. By default this information is added to the table, altering the stacked table schema from that of the raw table. Disable this behavior by setting `sensor_metadata = FALSE`. Future NEON sensor data products may start including this information in the raw data files, as is already the case for observational data.

## Examples

```
neon_read("brd_countdata-expanded")

## Sensor inputs will add metadata columns by default
neon_read("waq_instantaneous", site = c("CRAM", "SUGG"))
```

---

neon_remote	<i>neon_remote select a table from the remote connection</i>
-------------	--

---

**Description**

neon\_remote  
select a table from the remote connection

**Usage**

```
neon_remote(table = "", product = "", type = "", db = neon_remote_db())
```

**Arguments**

table	table name (pattern match regex)
product	product code
type	basic or extended (if necessary to distinguish)
db	a <a href="#">neon_remote_db</a> connection. If not provided, one will be created, but it is faster to pass this on for re-use in multiple neon_remote calls.

**Value**

a `arrow::FileSystemDataset` object, or a named list of such objects if multiple matches are found. This table is not downloaded but remains on the remote storage location, but can be filtered with `dplyr` functions like `filter` and `select`, and can also be grouped and summarised, all without ever downloading the whole table. Use `dplyr::collect()` to download the (possibly filtered) table into and pull into memory.

---

neon_remote_db	<i>Establish a remote database connection using arrow</i>
----------------	---

---

**Description**

Establish a remote database connection using arrow

**Usage**

```
neon_remote_db(
  bucket = arrow::s3_bucket("neon4cast-targets/neon", endpoint_override =
    "data.ecoforecast.org")
)
```

**Arguments**

bucket	an <code>[arrow::s3_bucket]</code> connection or other <code>arrow::SubTreeFileSystem</code> object.
--------	--

**Examples**

```
db <- neon_remote_db()
```

---

neon_sites	<i>Table of all NEON sites</i>
------------	--------------------------------

---

**Description**

Returns a table of all NEON sites by making a single API call to the /sites endpoint.

**Usage**

```
neon_sites(
  api = "https://data.neonscience.org/api/v0",
  .token = Sys.getenv("NEON_TOKEN")
)
```

**Arguments**

api	the URL to the NEON API, leave as default.
.token	an authentication token from NEON. A token is not required but will allow access to a higher number of requests before rate limiting applies, see <a href="https://data.neonscience.org/data-api/rate-limiting/#api-tokens">https://data.neonscience.org/data-api/rate-limiting/#api-tokens</a> . Note that once files are downloaded once, neonstore provides persistent access to them without further interaction required with the API.

---

neon_store	<i>import neon data into a local database</i>
------------	---

---

**Description**

import neon data into a local database

**Usage**

```
neon_store(
  table = NA,
  product = NA,
  type = NA,
  dir = neon_dir(),
  db = neon_db(neon_db_dir(), read_only = FALSE),
  n = 500L,
  quiet = FALSE,
  ...
)
```

**Arguments**

table	Include only files matching this table name (or regex pattern). (optional).
product	A NEON productCode or list of product codes, see examples.
type	Should we prefer the basic or expanded version of this product? Note that not all products have expanded formats.
dir	Location where files should be downloaded. By default will use the appropriate applications directory for your system (see <code>tools::R_user_dir()</code> ). This default also be configured by setting the environmental variable NEONSTORE_HOME, see <code>Sys.setenv</code> or <code>Renviron</code> .
db	A connection to a write-able relational database backend, see <code>neon_db()</code> .
n	number of files that should be read per iteration
quiet	show progress?
...	Arguments passed on to <code>neon_read</code>
sensor_metadata	logical, default TRUE. Should we add metadata fields from file names of sensor data into the table? Adds DomainID, SiteID, horizontalPosition, verticalPosition, and publicationDate. Results in slower parsing.
keep_filename	Should we include a column indicating the original file name for each row? Can be a useful source of additional metadata that NEON may omit from the raw files (i.e. siteID), but will also result in slower parsing. Default FALSE.
altrep	enable or disable altrep. Logical, default FALSE. Setting to TRUE can speed up reading, but may cause <code>vroom::vroom</code> to throw mapping error: Too many open files.
files	optionally, specify a vector of file paths directly (e.g. as provided from <code>neon_index</code> ) and specify table argument as NULL.
ext	only match files with this file extension(s)
timestamp	only match timestamps prior this. See details in <code>neon_index()</code> . Should be a datetime POSIXct object (or coerce-able string)
start_date	Download only files as recent as (YYYY-MM-DD). Leave as NA to download up to the most recent available data.
end_date	Download only files up to end_date (YYYY-MM-DD). Leave as NA to download all prior data.
site	4-letter site code(s) to filter on. Leave as NA to search all.
release	Select only data files associated with a particular release tag, see <a href="https://www.neonscience.org/data-samples/data-management/data-revisions-releases">https://www.neonscience.org/data-samples/data-management/data-revisions-releases</a> , e.g. "RELEASE-2021". Releases are associated with a specific DOI and the promise that files associated with a particular release will not change.

**Value**

the index of files read in (invisibly)



---

neon_sync_db	<i>sync local parquet export to an S3 database</i>
--------------	--

---

**Description**

sync local parquet export to an S3 database

**Usage**

```
neon_sync_db(s3, dir = file.path(neon_dir(), "parquet"))
```

**Arguments**

s3	an [arrow::SubTreeFileSystem], such as a remote connection to an S3 bucket from [arrow::s3_bucket()].
dir	directory to which parquet export is written.

**Details**

Remote files are named according to the table name (including product id, not according to the 'sanitized' file name duckdb uses when generating exports.)

---

neon_table	<i>Return a neon table from the database</i>
------------	--

---

**Description**

Return a neon table from the database

**Usage**

```
neon_table(  
  table,  
  product = NA,  
  type = NA,  
  site = NA,  
  db = neon_db(),  
  lazy = FALSE  
)
```

**Arguments**

table	the name of a downloaded NEON table in the store, see <a href="#">neon_index</a>
product	A NEON productCode or list of product codes, see examples.
type	filter for basic or expanded. Can be omitted unless you have imported both types a given table into your database.
site	4-letter site code(s) to filter on. Leave as NA to search all.
db	a connection to the database, see <code>[neon_db()]</code> .
lazy	logical, default FALSE. Should we return a remote dplyr connection to the table in duckdb? This can substantially improve performance and avoid out-of-memory errors when working with very large tables. However, not all R operations can be performed on a remote table, only (most) functions from dplyr and tidyr, as these can be translated automatically to SQL language used by the remote database. Use dplyr functions like <code>dplyr::filter()</code> , <code>dplyr::group_by()</code> , and <code>dplyr::summarise()</code> to subset the data appropriately within the remote table before calling <code>[dplyr::collect()]</code> to import the data fully into R.

**Details**

We cannot filter on `start_date` or `end_date` since these come only from the filename metadata and are only added to instrument tables, not observation tables etc.

---

`show_deprecated_data` *show deprecated data*

---

**Description**

Show the file information for any raw data files which have been deprecated by the release of modified historical data to the NEON API.

**Usage**

```
show_deprecated_data(
  product = NA,
  table = NA,
  site = NA,
  start_date = NA,
  end_date = NA,
  type = NA,
  ext = NA,
  timestamp = NA,
  release = NA,
  dir = neon_dir()
)
```

**Arguments**

product	A NEON productCode or list of product codes, see examples.
table	Include only files matching this table name (or regex pattern). (optional).
site	4-letter site code(s) to filter on. Leave as NA to search all.
start_date	Download only files as recent as (YYYY-MM-DD). Leave as NA to download up to the most recent available data.
end_date	Download only files up to end_date (YYYY-MM-DD). Leave as NA to download all prior data.
type	Should we prefer the basic or expanded version of this product? Note that not all products have expanded formats.
ext	only match files with this file extension(s)
timestamp	only match timestamps prior this. See details in <a href="#">neon_index()</a> . Should be a datetime POSIXct object (or coerce-able string)
release	Select only data files associated with a particular release tag, see <a href="https://www.neonscience.org/data-samples/data-management/data-revisions-releases">https://www.neonscience.org/data-samples/data-management/data-revisions-releases</a> , e.g. "RELEASE-2021". Releases are associated with a specific DOI and the promise that files associated with a particular release will not change.
dir	Location where files should be downloaded. By default will use the appropriate applications directory for your system (see <a href="#">tools::R_user_dir()</a> ). This default also be configured by setting the environmental variable NEONSTORE_HOME, see <a href="#">Sys.setenv</a> or <a href="#">Renviron</a> .

**Details**

NEON data files are sometimes updated to correct errors. Old files are removed from access from the API, but may be present in your local store from an earlier download. `neonstore` stacking functions (`[neon_read()]` and `neon_store()`) automatically exclude these deprecated files, though `neon_read()` can be instructed to use older files by passing a file list.

A data file is identified as deprecated whenever the local file store contains a second data file with the same product, table, site, month, and position (sensor products only) information, but having an updated timestamp. If such a change occurs in a file with a non-missing "month" code, it may indicate a data file has been updated. This could result in changes to the results of any previous analyses.

Note that metadata files, (readme, variables, positions) are 'pre-stacked': the metadata file in a given product-site-month set contains metadata going back to the start and not just for that month. As a result, each new version deprecates the old metadata file, but the old files are always available from the NEON API and always present in the store. Users will only need to care about the most recent ones, and the presence of old files is no cause for concern. This function will only show data files that have changed, and not metadata files. This can help pinpoint specific altered data.

**See Also**

`neon_index`, `neon_read`

**Examples**

```
show_deprecated_data()
```

---

```
standardize_export_names
```

*standardize export names*

---

**Description**

standardize export names

**Usage**

```
standardize_export_names(dir = file.path(neon_dir(), "parquet"))
```

**Arguments**

`dir`                    directory to which parquet export is written.

**Details**

DUCKDB clobbers database filenames to avoid potentially incompatible characters. This is pretty unnecessary, so we can restore the original table names for use with S3-based remote access which assumes parquet files map to the desired table names (i.e. including product numbers.)

However, note that `[neon_import_db()]` uses native duckdb functions that assume the original mangled names.

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