

# Package: wcUtils (via r-universe)

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**Title** Utilities to Access the Wildlife Computers Data Portal API

**Version** 0.1

**Description** This package provides basic access and capabilities for interacting with and downloading data from the Wildlife Computers Data Portal API. Where appropriate functions to specify data types and tidy the data are employed.

**Depends** R (>= 3.1.1)

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**Repository** <https://ocean-tracking-network.r-universe.dev>

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as\_ecdf

*Create An ECDF List*


---

## Description

This creates a standardized list for holding ECDF data and sets the class to ‘ecdf’. The S3 class, \*wcECDF\* consists of three named objects: ‘type’, ‘percent\_time’, and ‘ecdf’. The ‘type’ object is the same as that specified for the ‘type’ parameter. The ‘percent\_time’ object holds the proportion of the summary period the tag spent within the ‘shallow’ or ‘deep’ portion of the water column. Lastly, the ‘ecdf’ object is a data.frame with a column, ‘ecd\_prop’, that specifies the proportional values for the ECDF and a column, ‘depth\_break’, that reports the corresponding depth value (in meters).

## Usage

```
as_ecdf(type, ecdf)
```

## Arguments

|      |                                    |
|------|------------------------------------|
| type | this is either ‘shallow’ or ‘deep’ |
| ecdf | a data.frame                       |

## Value

A list with three named objects (‘type’, ‘percent\_time’, ‘ecdf’) of class ‘wcECDF’

---

|              |  |
|--------------|--|
| combine_ecdf | <i>Combine Two 'wcECDF' Objects Into a Single ECDF</i> |
|--------------|--|

---

**Description**

Combine Two 'wcECDF' Objects Into a Single ECDF

**Usage**

```
combine_ecdf(ecdf1, ecdf2, return = "pdf")
```

**Arguments**

|       |   |
|-------|---|
| ecdf1 | object of class 'wcECDF'; typically the <i>*shallow*</i> region |
| ecdf2 | object of class 'wcECDF'; typically the <i>*deep*</i> region    |

**Value**

four-column data.frame with columns 'ecd\_prop', 'pdf', 'prob', and 'depth\_break'

---

|                     |                                       |
|---------------------|---------------------------------------|
| drop_ecd_duplicates | <i>Drop duplicates within an ECDF</i> |
|---------------------|---------------------------------------|

---

**Description**

Drop duplicates within an ECDF

**Usage**

```
drop_ecd_duplicates(e)
```

**Arguments**

e

---

 fixCSV

*Tidy a comma separated value (CSV) file*


---

### Description

Tidies up a Comma Separated Value (CSV) file, ensuring that each row of the table in the file contains the same number of commas, and no empty rows are left below the table.

### Usage

```
fixCSV(file, skip = 0, overwrite = FALSE)
```

### Arguments

|           |  |
|-----------|--|
| file      | character: the name of the CSV file to be ‘fixed’.   |
| skip      | integer: the number of lines in the CSV file to skip before the header row of the table. The skipped lines are copied directly to the output file unchanged. The default is skip=0, implying that the header row is the first row of the CSV file. |
| overwrite | logical: Write output to a separate, ‘FIXED’ file (overwrite=FALSE, the default), or overwrite the original file (overwrite=TRUE)? If overwrite=TRUE, the original file is copied to a .BAK file before being overwritten.                         |

### Details

fixCSV tidies up a Comma Separated Value (CSV) file to ensure that the CSV file contains a strictly rectangular block of data for input into R (ignoring any preliminary comment rows via the skip= argument).

CSV formatted files are a plain text file format for tabular data, in which cell entries in the same row of a table are separated by commas. When such files are exported from other applications such as spreadsheet software, the software has to decide whether any empty cells to the right-hand side of, or below, the table or spreadsheet should be represented by trailing commas in the CSV file. Such decisions can result in a ‘ragged’ table in the CSV file, in which some rows contain fewer commas (‘short rows’) or more commas (‘long rows’) than others, or where empty rows below the table are included as comma-only rows in the CSV file.

While R’s [read.table](#) and related functions can sensibly extend short rows as needed, ragged tables in a CSV file can still result in errors, unwanted empty rows (below the table) or unwanted columns (to the right of the table) when the data is loaded into R.

fixCSV reads in a specified CSV file and removes or adds commas to rows, to ensure that each row in the body of the table contains the same number of cells as the header row of the table. Any empty rows below the table are also removed. The resulting table is then written back to file, either to a new file with ‘FIXED’ added to the filename (argument overwrite=FALSE, the default) or overwriting the original file (overwrite=TRUE - the original file is copied to a .BAK file before being overwritten).

Note that:

- The table of data in the CSV file *must* contain a header row of the correct length, since this row is used to determine the correct number of columns for the table. Note: if this header row is too short, then subsequent rows will be truncated to match the length of the header, so beware. Misspecification of the skip= argument (see below) can similarly lead to such corruption of the 'fixed' file.
- In the header row, any trailing commas representing empty cells to the right of the (non-empty) header entries are first removed before determining the correct number of columns for the table. Thus the length of the header row (and hence the assumed width of the entire table) is determined by the *right-most non-empty cell* in the header row.
- fixCSV does not remove empty cells, rows or columns within the interior (or on the left side) of the table - it is concerned only with the right and bottom boundaries of the table.
- A skip= argument is included to tell fixCSV to ignore the specified number of comment rows preceding the header row. Such rows are simply copied over into the output file unchanged. The default for this parameter is skip=0, so that the first row in the data file is assumed to be the header row. As noted above, misspecification of this argument can seriously corrupt the output.
- fixCSV can overwrite your data file(s) (via overwrite=TRUE), and although it makes a backup of your original file, you should still make sure that you have a separate backup of your data file in a safe place before using this function! The author of this code takes no responsibility for any data loss or corruption as a result of the use of this routine...

### Author(s)

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### Examples

```
## Not run:  
  
## Assuming CSV file 'alleleDataFile.csv' exists in the current  
## directory. The following overwrites the CSV file - make sure  
## you have a backup!  
  
fixCSV("alleleDataFile.csv", overwrite=TRUE)  
  
## End(Not run)
```

---

format\_bins

*format bin labels*

---

### Description

format bin labels

**Usage**

```
format_bins(bins)
```

**Arguments**

bins                    a vector of bin labels to be formatted

**Value**

a vector of formatted bin labels

---

|            |  |
|------------|--|
| make_names | <i>Expanded make.names function for creating consistent column names</i> |
|------------|--|

---

**Description**

Expanded make.names function for creating consistent column names

**Usage**

```
make_names(x)
```

**Arguments**

x                        data frame with columns to be renamed

**Value**

a data frame

---

|             |  |
|-------------|--|
| read_allmsg | <i>Parse a *-All.csv file into a proper data.frame</i> |
|-------------|--|

---

**Description**

Parse a \*-All.csv file into a proper data.frame

**Usage**

```
read_allmsg(allmsg_file, to_lower = TRUE, fix_csv = FALSE)
```

**Arguments**

allmsg\_file            file path or file connection to a \*-All.csv file  
to\_lower                whether to convert the column names to lower case  
fix\_csv                 whether to attempt to fix any comma, csv issues

**Value**

a data frame

---

|            |   |
|------------|---|
| read_behav | <i>Parse a *-Behavior.csv file into a proper data.frame</i> |
|------------|---|

---

**Description**

Parse a \*-Behavior.csv file into a proper data.frame

**Usage**

```
read_behav(behav_file, to_lower = TRUE, fix_csv = FALSE)
```

**Arguments**

|            |   |
|------------|---|
| behav_file | file path or file connection to a *-Behavior.csv file |
| to_lower   | whether to convert the column names to lower case     |
| fix_csv    | whether to attempt to fix any comma, csv issues       |

**Value**

a data frame

---

|           |                                     |
|-----------|-------------------------------------|
| read_ecdf | <i>Read and Tidy ECDF Data File</i> |
|-----------|-------------------------------------|

---

**Description**

This function is the workhorse of the 'wcECDF' package. All ECDF data are stored within a '\*-ECDHistos.csv' file that is output from either the Wildlife Computers Data Portal or DAP processing software. The function presumes the '\*-ECDHistos.csv' data file is provided as-is from these sources and has not been edited. The resulting output is a nested 'tibble' that adheres to tidy data principles and includes new columns ('shallow\_ecdf', 'deep\_ecdf', 'full\_ecdf', and 'full\_pdf').

**Usage**

```
read_ecdf(ecdf_csv)
```

**Arguments**

|          |                                     |
|----------|-------------------------------------|
| ecdf_csv | file path for the '*-ECDHistos.csv' |
|----------|-------------------------------------|

## Details

In addition to *tidying* up the original data into a more workable *long* format, this function calculates four new columns.

**shallow\_ecdf** The `shallow_ecdf` column is a list-col that contains nested S3 objects of class `wcECDF` representing the portion of the water column defined as `shallow`.

**deep\_ecdf** The `deep_ecdf` column is a list-col that contains nested S3 objects of class `wcECDF` representing the portion of the water column defined as `deep`.

**full\_ecdf** The combined ECDF for both shallow and deep regions. The resulting ECDF is weighted based on the reported proportion of time spent within each region.

**full\_pdf** The `full_ecdf` is transformed into a probability density function and two columns are returned: `pdf` and `prob`. The later represents the probability the tag spent time at a given depth.

## Value

A nested tibble

---

|              |   |
|--------------|---|
| read_fastGPS | <i>Parse a *-FastGPS.csv files into a proper data.frame</i> |
|--------------|---|

---

## Description

Parse a *-FastGPS.csv* files into a proper `data.frame`

## Usage

```
read_fastGPS(gps_file, to_lower = TRUE, fix_csv = FALSE)
```

## Arguments

|                       |  |
|-----------------------|--|
| <code>gps_file</code> | file path or file connection to a <i>-FastGPS.csv</i> file |
| <code>to_lower</code> | whether to convert the column names to lower case          |
| <code>fix_csv</code>  | whether to attempt to fix any comma, csv issues            |

## Value

a data frame



---

read\_histos                      *Parse a \*-Histos.csv files into a proper data.frame*

---

**Description**

Parse a \*-Histos.csv files into a proper data.frame

**Usage**

```
read_histos(  
  histo_file,  
  to_lower = TRUE,  
  dt_fmt = "%H:%M:%S %d-%b-%Y",  
  fix_csv = FALSE  
)
```

**Arguments**

|            |   |
|------------|---|
| histo_file | file path or file connection to a *-Histos.csv file |
| to_lower   | whether to convert the column names to lower case   |
| dt_fmt     | format for the Date column                          |
| fix_csv    | whether to attempt to fix any comma, csv issues     |

**Value**

a list of two data frames

---

read\_locs                      *Parse a \*-Locations.csv files into a proper data.frame*

---

**Description**

Parse a \*-Locations.csv files into a proper data.frame

**Usage**

```
read_locs(loc_file, fix_csv = FALSE)
```

**Arguments**

|          |  |
|----------|--|
| loc_file | file path or file connection to a *-Locations.csv file |
| fix_csv  | whether to attempt to fix any comma, csv issues        |

**Value**

a data frame

---

|             |                             |
|-------------|-----------------------------|
| smooth_ecdf | <i>Create smoothed ECDF</i> |
|-------------|-----------------------------|

---

**Description**

Create smoothed ECDF

**Usage**

```
smooth_ecdf(ecdf, bin.width)
```

**Arguments**

bin.width

---

|                |   |
|----------------|---|
| tidyDiveDepths | <i>Apply 'tidy' data principles to the dive-depth histogram data stream</i> |
|----------------|---|

---

**Description**

tidyDiveDepths returns a 'tidy'd' data frame of dive depth data

**Usage**

```
tidyDiveDepths(histos)
```

**Arguments**

histos            a list returned from read\_histos

**Details**

The histogram data stream is provided in a 'wide' format (each row represents a time period and the observed values are provided in 1 to 72 'bin' columns). This format can be difficult to work with in R and other data analysis platforms (e.g. database tables), so we use the `tidyr` and `dplyr` packages to manipulate the data into a more flexible, 'narrow' format. This results in a data structure where every row represents a single observation.

This is implemented, here, with the dive depth data. For dive depth data, the tag records the maximum depth experienced during a qualifying dive and tallies those dives into user-specified depth bins and user-specified time bins. This, unlike with timeline data, requires some knowledge of these user-specified bins. As long as the user has uploaded a configuration/report file to the Wildlife Computers Data Portal, then the `*-Histos.csv` file provides information on the dive depth bins. If the bin information is not available, the function will produce a warning and output files with generic 'Bin' labels.

**Value**

a data frame with tidy, narrow data structure and actual dive depths bin limits (when provided)

---

|                   |  |
|-------------------|--|
| tidyDiveDurations | <i>Apply 'tidy' data principles to the dive-duration histogram data stream</i> |
|-------------------|--|

---

**Description**

tidyDiveDurations returns a 'tidy'd' data frame of dive duration data

**Usage**

```
tidyDiveDurations(histos)
```

**Arguments**

histos            a list returned from read\_histos

**Details**

The histogram data stream is provided in a 'wide' format (each row represents a time period and the observed values are provided in 1 to 72 'bin' columns). This format can be difficult to work with in R and other data analysis platforms (e.g. database tables), so we use the `tidyr` and `dplyr` packages to manipulate the data into a more flexible, 'narrow' format. This results in a data structure where every row represents a single observation.

This is implemented, here, with the dive duration data. For dive duration data, the tag records the duration in seconds of a qualifying dive and tallies those durations into user-specified duration bins and user-specified time bins. This, unlike with timeline data, requires some knowledge of these user-specified bins. As long as the user has uploaded a configuration/report file to the Wildlife Computers Data Portal, then the \*-Histos.csv file provides information on the dive durations. If the bin information is not available, the function will produce a warning and output files with generic 'Bin' labels.

**Value**

a data frame with tidy, narrow data structure and actual dive duration bin limits (when provided)

---

|                 |  |
|-----------------|--|
| tidyTimeAtDepth | <i>Apply 'tidy' data principles to the time-at-depth histogram data stream</i> |
|-----------------|--|

---

**Description**

tidyTimeAtDepth returns a 'tidy'd' data frame of time-at-depth data

**Usage**

```
tidyTimeAtDepth(histos)
```

**Arguments**

histos            a list returned from read\_histos

**Details**

The histogram data stream is provided in a 'wide' format (each row represents a time period and the observed values are provided in 1 to 72 'bin' columns). This format can be difficult to work with in R and other data analysis platforms (e.g. database tables), so we use the `tidyr` and `dplyr` packages to manipulate the data into a more flexible, 'narrow' format. This results in a data structure where every row represents a single observation.

This is implemented, here, with the time-at-depth data. For time-at-depth data, the tag records the portion of time the tag spent within user-defined dive depth bins. This, unlike with timeline data, requires some knowledge of these user-specified bins. As long as the user has uploaded a configuration/report file to the Wildlife Computers Data Portal, then the \*-Histos.csv file provides information on the time-at-depth bins. If the bin information is not available, the function will produce a warning and output files with generic 'Bin' labels.

**Value**

a tibble with tidy, narrow data structure and actual time-at-depth bin limits (when provided)

---

|               |   |
|---------------|---|
| tidyTimelines | <i>Apply 'tidy' data principles to the timeline histogram data stream</i> |
|---------------|---|

---

**Description**

tidyTimelines returns a 'tidy'd' data frame of timeline data

**Usage**

```
tidyTimelines(histos, row_min = 1)
```

**Arguments**

|         |   |
|---------|---|
| histos  | a list returned from read_histos  |
| row_min | user defined minimum timeline records required; row_min = 2 will return NULL unless at least 1 records are found. |

**Details**

The histogram data stream is provided in a 'wide' format (each row represents a time period and the observed values are provided in 1 to 72 'bin' columns). This format can be difficult to work with in R and other data analysis platforms (e.g. database tables), so we use the `tidyr` and `dplyr` packages to manipulate the data into a more flexible, 'narrow' format. This results in a data structure where every row represents a single observation.

This is implemented, here, with the timeline data. For timeline data, tag 'dryness' is provided as either a percentage of each hour the tag was dry or as a binary (1 or 0) value representing whether a tag was dry for a majority of a given 20-minute period. For both of these situations, the values for the 'bin' columns are predictable and we can, in addition to tidying the data structure, also turn the bin values into actual time periods.

**Value**

a data frame with tidy, narrow data structure and actual time periods in place of bins

---

|        |  |
|--------|--|
| to_pdf | <i>Transform a 'weECDF' Object Into a Probability Density Function</i> |
|--------|--|

---

**Description**

Transform a 'weECDF' Object Into a Probability Density Function

**Usage**

```
to_pdf(ecdf)
```

**Arguments**

|      |  |
|------|--|
| ecdf | two-column data.frame with 'ecdf_prop' and 'depth_break' |
|------|--|

**Value**

a two-column data.frame with 'ecdf\_prop' and 'depth\_break'

---

|               |   |
|---------------|---|
| wcGetDeployID | <i>Return a vector of data portal unique IDs associated with a given DeployID</i> |
|---------------|---|

---

### Description

wcGetDeployID returns a vector of data portal unique ID(s)

### Usage

```
wcGetDeployID(xml_content, deployid = NULL)
```

### Arguments

|             |   |
|-------------|---|
| xml_content | XML content/data returned from wcPOST (with 'action=get_deployments') |
| deployid    | valid deployid character (required)                                   |

### Details

This function presumes a DeployID has been setup for deployments on the Wildlife Computers Data Portal. The vector of deployment ID(s) returned will be a subset that match the deployid character in the function call. The list returned will also include a simple data frame with summary information one can use to determine the appropriate id.

### Value

a list with ids (a vector of deployment ids) and a df (data frame of deployment summaries)

---

|               |  |
|---------------|--|
| wcGetDownload | <i>Retrieve data from Wildlife Computers Data Portal</i> |
|---------------|--|

---

### Description

wcGetDownload will return a list of data frames containing deployment data

### Usage

```
wcGetDownload(
  id,
  wc.key = Sys.getenv("WCACCESSKEY"),
  wc.secret = Sys.getenv("WCSECRETKEY"),
  keyfile = NULL,
  tidy = TRUE
)
```

**Arguments**

|      |  |
|------|--|
| id   | a single character representing a data portal unique deployment identifier |
| tidy | whether to tidy the histogram data stream and create a timelines output    |

**Details**

The Wildlife Computers Data Portal will return deployment data in the form of a zipped file with various comma-separated files and other accessory files. The \*.csv files correspond to particular data streams. This function, currently, focuses on the locations, behavior, histograms, timelines, status, and messages data streams.

For most of the files, the data are read in with `read.csv` and, other than a few steps to set the data types, the data are provided 'as is'. The one exception is the histogram data stream. Here, we use the `tidyr` and `dplyr` package to 'tidy' the data into a more appropriate data structure. For now, this is only implemented with timeline data and the 'tidy'd' data is provided within the list element `$timelines`

**Value**

a list of data frames with up to 6 named elements (locations, behavior, histograms, status, timelines, messages)

---

 wcGetIDs

*Return a vector of deployment IDs*


---

**Description**

wcGetIDs returns a vector of deployment IDs

**Usage**

```
wcGetIDs(xml_content, xpath = NULL)
```

**Arguments**

|             |   |
|-------------|---|
| xml_content | XML content/data returned from wcPOST (with 'action=get_deployments') |
| xpath       | additional customization possible by passing an xpath statement       |

**Details**

Each deployment in the Wildlife Computers Data Portal is identified by a unique alpha-numeric value. This function searches the XML response data and extracts those IDs

**Value**

returns a vector of deployment IDs

---

wcGetProjectIDs      *Return a vector of deployment IDs associated with a given Project*

---

### Description

wcGetProjectIDs returns a vector of deployment IDs

### Usage

```
wcGetProjectIDs(xml_content, project = NULL)
```

### Arguments

|             |   |
|-------------|---|
| xml_content | XML content/data returned from wcPOST (with 'action=get_deployments') |
| project     | valid project name (required)   |

### Details

This function presumes a custom label, 'Project', has been setup for deployments on the Wildlife Computers Data Portal. The vector of deployment IDs returned will be a subset that match the project name provided in the function call.

### Value

returns a vector of deployment IDs

---

wcGetPttID      *Return a vector of deployment ID associated with a given PTT*

---

### Description

wcGetPttID returns a vector of deployment ID(s)

### Usage

```
wcGetPttID(xml_content, ptt = NULL)
```

### Arguments

|             |   |
|-------------|---|
| xml_content | XML content/data returned from wcPOST (with 'action=get_deployments') |
| ptt         | valid ptt integer (required)  |



**Details**

This function presumes a PTT has been setup for deployments on the Wildlife Computers Data Portal. The vector of deployment ID(s) returned will be a subset that match the ptt integer provided in the function call. The list returned will also include a simple data frame with summary information one can use to determine the appropriate id.

**Value**

a list with ids (a vector of deployment ids) and a df (data frame of deployment summaries)

---

|                |   |
|----------------|---|
| wcGetRecentIDs | <i>Return a vector of deployment IDs with new data in the last n days</i> |
|----------------|---|

---

**Description**

wcGetProjectIDs returns a vector of deployment IDs

**Usage**

```
wcGetRecentIDs(xml_content, days = 14)
```

**Arguments**

|             |   |
|-------------|---|
| xml_content | XML content/data returned from wcPOST (with 'action=get_deployments') |
| days        | integer value specifying the time window from now() in days           |

**Details**

This returns a subset of deployment IDs with new data available on the portal within the last n days. The default value is for 14 days

**Value**

returns a vector of deployment IDs

wcGetZip

*Retrieve a single zip file from Wildlife Computers Data Portal*

---

**Description**

wcGetZip will return a path to a downloaded zip file

**Usage**

```
wcGetZip(  
    id,  
    wc.key = Sys.getenv("WCACCESSKEY"),  
    wc.secret = Sys.getenv("WCACCESSKEY"),  
    keyfile = NULL  
)
```

**Arguments**

|           |  |
|-----------|--|
| wc.key    | public access key (default retrieves from option value set in .Renviron) |
| wc.secret | secret access key (default retrieves from option value set in .Renviron) |
| keyfile   | path to a json formatted keyfile with WCACCESSKEY and wcSecretKey        |

**Details**

The Wildlife Computers Data Portal will return deployment data in the form of a zipped file with various comma-separated files and other accessory files.

**Value**

a path to the zip file

---

wcPOST

*Send POST to Wildlife Computers Data Portal API*

---

**Description**

wcPOST returns a response from a POST to the API

**Usage**

```
wcPOST(  
    wc.key = Sys.getenv("WCACCESSKEY"),  
    wc.secret = Sys.getenv("WCSECRETKEY"),  
    keyfile = NULL,  
    params = "action=get_deployments"  
)
```

**Arguments**

|           |  |
|-----------|--|
| wc.key    | public access key (default retrieves from option value set in .Renviron) |
| wc.secret | secret access key (default retrieves from option value set in .Renviron) |
| keyfile   | path to a json formatted keyfile with WCACCESSKEY and WCSECRETKEY        |
| params    | POST message (default returns a list of deployments)                     |

**Details**

This function provides basic access to the API via POST. The params value contains the string to include in the body of the POST. The default action is to return a list of deployments associated with your account. Most users will likely not call this function directly, but instead, rely on other helper/wrapper functions within the package.

The Wildlife Computers Data Portal API uses a form of keyed-hash message authentication code for secure access. Your 'Access Key' and 'Secret Key' can be obtained from the data portal website (Account Settings > Web Services Security). For security reasons you should NOT include the keys as plain text in any scripts. Instead, include the key values as within your .Renviron.

**Value**

an http response object is returned. Content of the response can be obtained with the http::content() function.

**Setting key values within .Renviron**

This option is a preferred option for storing keys, passwords and other sensitive values. Your .Renviron should be secured via OS security/permissions (e.g. on Linux/OS X .Renviron is stored within the home directory which is only accessible by an authorized user. However, you should not share your .Renviron or include your .Renviron in version control (e.g. git) if you use this option. An alternative is to read values from a different file in the home directory or to use OS level environment variables.

```
preformatted WCACCESSKEY = 'E4iZhsfdje7590JDNR/VARTEZyhfbw84485X5Xw86ow=' WC-
SECRETKEY = 'WIRJFYhfjdsuSEqKoE7WSDvXUHVP0pHDJSscmeA7fw='
```

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 wcUtils

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*wcUtils: A package for import/export of Wildlife Computers tag data*


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**Description**

wcUtils provides functionality for working with data from select (SPLASH, SPOT) Wildlife Computers satellite telemetry tags. The package relies on data files produced by the Wildlife Computers DAP program or downloaded from the Wildlife Computers Data Portal. Data are read into R

**wcUtils functions**

- read\_behav
- read\_histos
- read\_locs
- tidyDiveDepths
- tidyDiveDurations
- tidyTimeAtDepth
- tidyTimelines
- wcGetDownload
- wcGetIDs
- wcGetProjectIDs
- wcGetRecentIDs
- wcPOST

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