

# Package ‘vpr’

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**Type** Package

**Title** Processing and Visualization of Video Plankton Recorder Data

**Version** 0.2.3

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**Description** An oceanographic data processing package for analyzing and visualizing Video Plankton Recorder data. This package was developed at 'Bedford Institute of Oceanography'. Functions are designed to process automated image classification output and create organized and easily portable data products.

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**Encoding** UTF-8

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**BuildVignettes** true

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`bin_calculate`      *Get bin averages for VPR and CTD data*

---

**Description**

Bins CTD data for an individual cast to avoid depth averaging across tow-yo's

**Usage**

```
bin_calculate(data, binSize = 1, imageVolume, rev = FALSE)
```

**Arguments**

<code>data</code>	ctd data frame object including scan, salinity, temperature, depth, conductivity, time, fluor_ref, turbidity_ref, turbidity_mv, altitude, cast_id, n_roi
<code>binSize</code>	the height of bins over which to average, default is 1 metre
<code>imageVolume</code>	the volume of VPR images used for calculating concentrations (mm <sup>3</sup> )
<code>rev</code>	logical value, if TRUE, binning will begin at bottom of each cast, this controls data loss due to uneven binning over depth. If bins begin at bottom, small amounts of data may be lost at the surface of each cast, if binning begins at surface (rev = FALSE), small amounts of data may be lost at bottom of each cast

**Details**

Image volume calculations can change based on optical setting of VPR as well as autodeck setting used to process images For IML2018051 (S2) image volume was calculated as 108155 mm<sup>3</sup> by seascan (6.6 cubic inches) For COR2019002 S2 image volume was calculated as 83663 mm<sup>3</sup> and S3 image volume was calculated as 366082 mm<sup>3</sup>. Used internally ( `bin_cast` ) after `ctd_cast` on a single ascending or descending section of VPR cast

**Note**

`binSize` should be carefully considered for best results  
Depth is used for calculations! Please ensure depth is included in data frame using `swDepth`

**Author(s)**

E. Chisholm, K. Sorochan

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bin_cast	<i>Bin vpr data</i>
----------	---------------------

---

### Description

Formats oce style VPR data into depth averaged bins using [ctd\\_cast](#) and [bin\\_calculate](#) This function is used inside [concentration\\_category](#)

### Usage

```
bin_cast(ctd_roi_oce, imageVolume, binSize, rev = FALSE)
```

### Arguments

ctd_roi_oce	oce ctd format VPR data from <a href="#">vpr_oce_create</a>
imageVolume	the volume of VPR images used for calculating concentrations (mm <sup>3</sup> )
binSize	passed to <a href="#">bin_calculate</a> , determines size of depth bins over which data is averaged
rev	logical value,passed to <a href="#">bin_calculate</a> if TRUE, binning will begin at bottom of each cast, this controls data loss due to uneven binning over depth. If bins begin at bottom, small amounts of data may be lost at the surface of each cast, if binning begins at surface (rev = FALSE), small amounts of data may be lost at bottom of each cast

### Details

Image volume calculations can change based on optical setting of VPR as well as autodeck setting used to process images For IML2018051 (S2) image volume was calculated as 108155 mm<sup>3</sup> by seascan (6.6 cubic inches) For COR2019002 S2 image volume was calculated as 83663 mm<sup>3</sup> and S3 image volume was calculated as 366082 mm<sup>3</sup>

### Value

A dataframe of depth averaged bins of VPR data over an entire cast with calculated concentration values

---

concentration_category	<i>Binned concentrations</i>
------------------------	------------------------------

---

### Description

This function produces depth binned concentrations for a specified taxa. Similar to [bin\\_cast](#) but calculates concentrations for only one taxa. Used inside [vpr\\_roi\\_concentration](#)

**Usage**

```
concentration_category(data, taxa, binSize, imageVolume, rev = FALSE)
```

**Arguments**

data	dataframe produced by processing internal to vpr_roi_concentration
taxa	name of taxa isolated
binSize	passed to <a href="#">bin_calculate</a> , determines size of depth bins over which data is averaged
imageVolume	the volume of VPR images used for calculating concentrations (mm <sup>3</sup> )
rev	Logical value defining direction of binning, FALSE - bins will be calculated from surface to bottom, TRUE- bins will be calculated bottom to surface

**Details**

Image volume calculations can change based on optical setting of VPR as well as autodeck setting used to process images For IML2018051 (S2) image volume was calculated as 108155 mm<sup>3</sup> by seascan (6.6 cubic inches) For COR2019002 S2 image volume was calculated as 83663 mm<sup>3</sup> and S3 image volume was calculated as 366082 mm<sup>3</sup>

**Author(s)**

E. Chisholm

---

ctd\_cast

*Isolate ascending or descending section of ctd cast*

---

**Description**

This is an internal step required to bin data

**Usage**

```
ctd_cast(
  data,
  cast_direction = "ascending",
  data_type,
  cutoff = 0.1,
  breaks = NULL
)
```

**Arguments**

data	an oce ctd object
cast_direction	'ascending' or 'descending' depending on desired section
data_type	specify 'oce' or 'df' depending on class of desired output
cutoff	Argument passed to <a href="#">ctdFindProfiles</a>
breaks	Argument passed to <a href="#">ctdFindProfiles</a>

**Value**

Outputs either data frame or oce ctd object

**Note**

[ctdFindProfiles](#) arguments for minLength and cutOff were updated to prevent losing data (EC 2019/07/23)

**Author(s)**

K Sorochan, E Chisholm

---

ctd_dat_combine	<i>VPR CTD data</i>
-----------------	---------------------

---

**Description**

A dataframe including all CTD parameters from the VPR CTD, produced by [vpr\\_ctd\\_read](#)

**Usage**

```
ctd_dat_combine
```

**Format**

A dataframe with 15 variables

**time\_ms** Time stamp when ROI was collected (milliseconds)

**conductivity** Conductivity collected by the VPR CTD

**pressure** Pressure measured from the VPR CTD in decibars

**temperature** Temperature measured from the VPR CTD in celsius

**salinity** Salinity measured from the VPR CTD

**fluor\_ref** A reference fluorescence baseline provided in millivolts by the VPR CTD for calibrating fluorescence\_mv data

**fluorescence\_mv** Fluorescence in millivolts from the VPR CTD (uncalibrated)

**turbidity\_ref** A reference turbidity baseline provided in millivolts for calibrating turbidity\_mv

**turbidity\_mv** Turbidity in millivolts from the VPR CTD (uncalibrated)  
**altitude\_NA** Altitude data from the VPR CTD  
**day** Day on which VPR data was collected (from AutoDeck)  
**hour** Hour during which VPR data was collected (from AutoDeck)  
**station** Station identifier provided during processing  
**sigmaT** Density calculated from temperature, pressure and salinity data  
**depth** Depth in metres calculated from pressure

---

ctd_df_cols	<i>Read CTD data (SBE49) and Fluorometer data from CTD- VPR package</i>
-------------	---

---

### Description

Internal use [vpr\\_ctd\\_read](#)

### Usage

```
ctd_df_cols(x, col_list)
```

### Arguments

x	full filename (ctd .dat file)
col_list	list of CTD data column names

### Details

**WARNING** This is hard coded to accept a specific order of CTD data columns. The names and values in these columns can change based on the specific instrument and should be updated before processing data from a new VPR.

Text file format .dat file Outputs ctd dataframe with variables time\_ms, conductivity, temperature, pressure, salinity, fluor\_ref, fluorescence\_mv, turbidity\_ref, turbidity\_mv, altitude\_NA

### Author(s)

K. Sorochan, E. Chisholm

---

 ctd\_roi\_merge

*VPR CTD data combined with tabulated ROIs*


---

### Description

A dataframe representing CTD data which has been merged with tabulated ROIs in each category, produced by [vpr\\_ctdroi\\_merge](#)

### Usage

```
ctd_roi_merge
```

### Format

A dataframe with 28 variables

**time\_ms** Time stamp when ROI was collected (milliseconds)

**conductivity** Conductivity collected by the VPR CTD

**pressure** Pressure measured from the VPR CTD in decibars

**temperature** Temperature measured from the VPR CTD in celsius

**salinity** Salinity measured from the VPR CTD

**fluor\_ref** A reference fluorescence baseline provided in millivolts by the VPR CTD for calibrating fluorescence\_mv data

**fluorescence\_mv** Fluorescence in millivolts from the VPR CTD (uncalibrated)

**turbidity\_ref** A reference turbidity baseline provided in millivolts for calibrating turbidity\_mv

**turbidity\_mv** Turbidity in millivolts from the VPR CTD (uncalibrated)

**altitude\_NA** Altitude data from the VPR CTD

**day** Day on which VPR data was collected (from AutoDeck)

**hour** Hour during which VPR data was collected (from AutoDeck)

**station** Station identifier provided during processing

**sigmaT** Density calculated from temperature, pressure and salinity data

**depth** Depth in metres calculated from pressure

**roi** ROI identification number

**categories** For each category name (eg. bad\_image\_blurry, Calanus, krill), there is a line in the dataframe representing the number of ROIs identified in this category

**n\_roi\_total** Total number of ROIs in all categories for each CTD data point



---

ctd_roi_oce	<i>VPR data including CTD and ROI information</i>
-------------	---

---

**Description**

An oce formatted CTD object with VPR CTD and ROI data from package example data set.

**Usage**

```
ctd_roi_oce
```

**Format**

An oce package format, a 'CTD' object with VPR CTD and ROI data (1000 data rows)

---

getRoiMeasurements	<i>THIS FUNCTION HAS BEEN DEPRECATED</i>
--------------------	--

---

**Description**

pull roi measurements from all taxa, all files

**Usage**

```
getRoiMeasurements(taxafolder, nchar_folder, unit = "mm", opticalSetting)
```

**Arguments**

taxafolder	path to taxa folder (base – autoid folder)
nchar_folder	number of characters in basepath
unit	unit data will be output in, 'mm' (default – millimetres) or 'px' (pixels)
opticalSetting	VPR optical setting determining conversion between pixels and millimetres (options are 'S0', 'S1', 'S2', or 'S3')

**Note**

This function is very finicky, easily broken because it relies on character string splitting. taxaFolder argument should not end in a backslash, please check output carefully to ensure taxa names or ROI numbers have been properly sub string'd

---

insertRow	<i>INTERNAL USE ONLY quick data frame function from github to insert row inside dat frame</i>
-----------	---

---

**Description**

INTERNAL USE ONLY quick data frame function from github to insert row inside dat frame

**Usage**

```
insertRow(existingDF, newrow, r)
```

**Arguments**

existingDF	data frame
newrow	new row of data
r	index of new row

---

isopycnal_calculate	<i>Get vector to draw isopycnal lines on TS plot Used internally to create TS plots</i>
---------------------	---

---

**Description**

Get vector to draw isopycnal lines on TS plot Used internally to create TS plots

**Usage**

```
isopycnal_calculate(sal, pot.temp, reference.p = 0)
```

**Arguments**

sal	salinity vector
pot.temp	temperature vector in deg C
reference.p	reference pressure for calculation, set to 0

**Note**

: modified from source:[https://github.com/Davidatlarge/ggTS/blob/master/ggTS\\_DK.R](https://github.com/Davidatlarge/ggTS/blob/master/ggTS_DK.R)

**Author(s)**

E. Chisholm

---

normalize_matrix	<i>Normalize a matrix</i>
------------------	---------------------------

---

**Description**

take each element of matrix divided by column total

**Usage**

```
normalize_matrix(mat)
```

**Arguments**

mat	a matrix to normalize
-----	-----------------------

**Details**

Make sure to remove total rows before using with VP data

**Note**

used internally for visualization of confusion matrices

---

px_to_mm	<i>Get conversion factor for pixels to mm for roi measurements</i>
----------	--

---

**Description**

Used internally

**Usage**

```
px_to_mm(x, opticalSetting)
```

**Arguments**

x	an aidmea data frame (standard) to be converted into mm from pixels
opticalSetting	the VPR setting determining the field of view and conversion factor between mm and pixels

**Details**

converts pixels to mm using conversion factor specific to optical setting

Options for opticalSetting are 'S0', 'S1', 'S2', or 'S3'

---

roimeas\_dat\_combine     *VPR measurement data calculated by Visual Plankton*

---

**Description**

A data frame of measurement information for each ROI in the sample data set including long axis length, perimeter and area, produced by [vpr\\_autoid\\_read](#)

**Usage**

roimeas\_dat\_combine

**Format**

A data frame with 12 variables

**roi** Unique ROI identifier - 10 digit

**taxa** Category in which ROI has been classified by Visual Plankton

**day\_hour** day and hour in which data was collected (from Autodeck)

**Perimeter** The perimeter of the ROI in millimeters

**Area** The area of the ROI in millimeters

**width1** Width at a first point of the ROI in millimetres (defined in more detail in VPR manual)

**width2** Width at a second point of the ROI in millimetres (defined in more detail in VPR manual)

**width3** Width at a third point of the ROI in millimetres (defined in more detail in VPR manual)

**short\_axis\_length** The length in millimeters of the ROI along the shorter axis

**long\_axis\_length** The length in millimeters of the ROI along the longer axis

**station** Station identifier provided in processing

**time\_ms** Time stamp when ROI was collected in milliseconds

---

roi\_dat\_combine     *VPR ROI data*

---

**Description**

A dataframe including VPR ROI data from the sample dataset, produced by [vpr\\_autoid\\_read](#)

**Usage**

roi\_dat\_combine

**Format**

A dataframe with 13 variables

**roi** Unique ROI identifier - 8 digit

**categories** For each category name (eg. bad\_image\_blurry, Calanus, krill), there is a line in the dataframe representing the number of ROIs identified in this category

**time\_ms** Time stamp when ROI was collected (milliseconds)

---

size\_df\_f

*VPR size information dataframe*

---

**Description**

A sample data frame of size information from Visual Plankton outputs, processed using [vpr\\_ctdroisize\\_merge](#)

**Usage**

size\_df\_f

**Format**

An object of class `data.frame` with 14 rows and 14 columns.

**Details**

@format A dataframe with 14 variables including

**frame\_ID** Unique identifier for each VPR frame

**pressure** Pressure measured from the VPR CTD in decibars

**temperature** Temperature measured from the VPR CTD in celsius

**salinity** Salinity measured from the VPR CTD

**sigmaT** Density calculated from temperature, salinity and pressure

**fluorescence\_mv** Fluorescence measured by the VPR CTD in millivolts (uncalibrated)

**turbidity\_mv** Turbidity measured by the VPR CTD in millivolts (uncalibrated)

**roi** Unique ROI identification number - 10 digits, 8 digit millisecond time stamp and two unique digits to denote multiple ROIs within a millisecond

**taxa** Category in which ROI has been classified by Visual Plankton

**day\_hour** Day and hour in which data was collected, from AutoDeck processing

**long\_axis\_length** The length of the longest axis of the ROI image, measured by Visual Plankton

**station** Station identifier provided during processing

**time\_ms** Time stamp when ROI was collected (milliseconds)

**roi\_ID** ROI identification number- 8 digit time stamp, without unique 2 digit ending

---

taxa_conc_n	<i>A binned data frame of concentration data per category</i>
-------------	---

---

### Description

A 'binned' dataframe from sample VPR data, including concentrations of each category, where each data point represents a 5 metre bin of averaged VPR data. Produced using [vpr\\_roi\\_concentration](#)

### Usage

taxa\_conc\_n

### Format

A dataframe with 21 variables

**depth** Depth calculated from pressure in metres

**min\_depth** The minimum depth of the bin in metres

**max\_depth** The maximum depth of the bin in metres

**depth\_diff** The difference between minimum and maximum bin depth in metres

**min\_time\_s** The minimum time in seconds of the bin

**max\_time\_s** The maximum time in seconds of the bin

**time\_diff\_s** The difference between minimum and maximum time in a bin, in seconds

**n\_roi\_bin** The number of ROI observations in a bin

**conc\_m3** The concentration of ROIs in a bin, calculated based on image volume and number of frames per bin

**temperature** Temperature measured from the VPR CTD in celsius (averaged within the bin)

**salinity** Salinity measured from the VPR CTD (averaged within the bin)

**density** sigma T density calculated from temperature, salinity and pressure (averaged within the bin)

**fluorescence** Fluorescence measured by the VPR CTD in millivolts (uncalibrated) (averaged within the bin)

**turbidity** Turbidity measured by the VPR CTD in millivolts (uncalibrated) (averaged within the bin)

**avg\_hr** The mean time in which bin data was collected, in hours

**n\_frames** The number of frames captured within a bin

**vol\_sampled\_bin\_m3** The volume of the bin sampled in metres cubed

**toyo** Identifier of the tow-yo section which bin is a part of, either ascending or descending, appended by a number

**max\_cast\_depth** The maximum depth of the entire VPR cast

**taxa** The category in which ROIs in bin have been classified by Visual Plankton

**station** Station identifier provided during processing

---

vpr\_autoid\_check      *Checks manually created aid files for errors*

---

### Description

Removes any empty aid files after manual reclassification, checks for tow numbers and other meta-data to match. Performs check to ensure measurement and ROI files are the same length

### Usage

```
vpr_autoid_check(basepath, cruise, del)
```

### Arguments

basepath	basepath to autoid folder eg. C:/data/CRUISENAME/autoid/
cruise	name of cruise which is being checked
del	Logical value, if TRUE, empty files will be deleted (see warning), if FALSE, files WILL NOT be deleted (they will be listed in output)

### Details

**WARNING:** This function will delete empty aid and aidmeas files, permanently changing your directory. Consider making a back up copy before running this function.

### Value

text file (saved in working directory) named CRUISENAME\_aid\_file\_check.txt

### Author(s)

E Chisholm

---

vpr\_autoid\_copy      *Copy VPR images into folders*

---

### Description

Organize VPR images into folders based on classifications provided by visual plankton

### Usage

```
vpr_autoid_copy(basepath, day, hour, classifier_type, classifier_name, taxa)
```

**Arguments**

basepath	A file path to your autoid folder where VP data is stored eg. "C:\data\cruise_XXXXXXXXXXXXXXXXXX\
day	character string representing numeric day of interest
hour	character string representing hour of interest
classifier_type	character string representing the type of classifier (either 'svm', 'nn' or 'dual') from Visual Plankton
classifier_name	character string representing name of Visual Plankton classifier
taxa	optional list of character strings if you wish to only copy images from specific classification groups

**Value**

organized file directory where VPR images are contained with folders, organized by day, hour and classification, inside your basepath/autoid folder

---

vpr\_autoid\_create      *Modifies aid and aid mea files based on manual reclassification*

---

**Description**

Modifies aid and aid mea files based on manual reclassification

**Usage**

```
vpr_autoid_create(reclassify, misclassified, basepath, day, hour, mea = TRUE)
```

**Arguments**

reclassify	list of reclassify files (output from vpr_manual_classification())
misclassified	list misclassify files (output from vpr_manual_classification())
basepath	base path to auto ID folder eg 'E:/autoID_EC_07032019/'
day	day identifier for relevant aid & aidmeas files
hour	hour identifier for relevant aid & aidmeas files
mea	logical indicating whether or not there are accompanying measurement files to be created

**examples:**

```
basepath <- 'E:/autoID_EC_07032019/' day <- '289' hr <- '08' day_hour_files
<- paste0('d', day, '.h', hr) misclassified <- list.files(day_hour_files, pattern
= 'misclassified_', full.names = TRUE) reclassify <- list.files(day_hour_files,
pattern = 'reclassify_', full.names = TRUE) vpr_autoid_create(reclassify, mis-
classified, basepath)
```

**Author(s)**

E. Chisholm



---

vpr\_autoid\_read      *Read VPR aid files*

---

### Description

Read aid text files containing ROI string information or measurement data and output as a dataframe

### Usage

```
vpr_autoid_read(  
  file_list_aid,  
  file_list_aidmeas,  
  export,  
  station_of_interest,  
  opticalSetting,  
  warn = TRUE  
)
```

### Arguments

**file\_list\_aid**      a list object of aid text files, containing roi strings. Output from matlab Visual Plankton software.

**file\_list\_aidmeas**      a list object of aidmea text files, containing ROI measurements. Output from matlab Visual Plankton software.

**export**      a character string specifying which type of data to output, either 'aid' (roi strings) or 'aidmeas' (measurement data)

**station\_of\_interest**      Station information to be added to ROI data output, use NA if irrelevant

**opticalSetting**      Optional argument specifying VPR optical setting. If provided will be used to convert size data into mm from pixels, if missing size data will be output in pixels

**warn**      Logical, FALSE silences size data unit warnings

### Details

Only outputs either ROI string information OR measurement data

### Note

Full paths to each file should be specified

### Author(s)

E. Chisholm & K. Sorochan

**Examples**

```

station_of_interest <- 'test'
dayhour <- c('d222.h03', 'd222.h04')

#' #VPR OPTICAL SETTING (S0, S1, S2 OR S3)
opticalSetting <- "S2"
imageVolume <- 83663 #mm^3

auto_id_folder <- system.file('extdata/COR2019002/autoid/', package = 'vpr', mustWork = TRUE)
auto_id_path <- list.files(paste0(auto_id_folder, "/"), full.names = TRUE)

#' # Path to aid for each taxa
aid_path <- paste0(auto_id_path, '/aid/')
# Path to mea for each taxa
aidmea_path <- paste0(auto_id_path, '/aidmea/')

# AUTO ID FILES
aid_file_list <- list()
aidmea_file_list <- list()
for (i in 1:length(dayhour)) {
  aid_file_list[[i]] <-
    list.files(aid_path, pattern = dayhour[[i]], full.names = TRUE)
  # SIZE DATA FILES
  aidmea_file_list[[i]] <-
    list.files(aidmea_path, pattern = dayhour[[i]], full.names = TRUE)
}

aid_file_list_all <- unlist(aid_file_list)
aidmea_file_list_all <- unlist(aidmea_file_list)

# ROIs
roi_dat_combine <-
  vpr_autoid_read(
    file_list_aid = aid_file_list_all,
    file_list_aidmeas = aidmea_file_list_all,
    export = 'aid',
    station_of_interest = station_of_interest,
    opticalSetting = opticalSetting,
    warn = FALSE
  )

# MEASUREMENTS
roimeas_dat_combine <-
  vpr_autoid_read(
    file_list_aid = aid_file_list_all,
    file_list_aidmeas = aidmea_file_list_all,
    export = 'aidmeas',
    station_of_interest = station_of_interest,
    opticalSetting = opticalSetting,
    warn = FALSE
  )

```

---

vpr_category	<i>Get taxa ids from string</i>
--------------	---------------------------------

---

**Description**

Get taxa ids from string

**Usage**

```
vpr_category(x)
```

**Arguments**

x                    A string specifying the directory of the "taxafolder", containing the taxa id

**Value**

A string of only the taxa id

**Author(s)**

K Sorochan

**See Also**

[vpr\\_hour](#), [vpr\\_day](#), [vpr\\_roi](#)

**Examples**

```
taxa_string <- 'C:/data/cruise/autoid/Calanus/d000/h00'  
vpr_category(taxa_string)
```

---

vpr\_category\_create    *Create a new taxa to be considered for classification after processing with VP*

---

### Description

creates empty directory structure to allow consideration of new taxa during vpr\_manual\_classification()

### Usage

```
vpr_category_create(taxa, basepath)
```

### Arguments

taxa	new taxa name to be added (can be a list of multiple taxa names)
basepath	basepath used for vpr_manual_classification

### Value

empty directory structure using new taxa name inside basepath

---

vpr\_ctdroisize\_merge    *Format CTD and Size data from VPR*

---

### Description

Format CTD and Meas data frames into combined data frame for analysis and plotting of size data

### Usage

```
vpr_ctdroisize_merge(data, data_mea, taxa_of_interest)
```

### Arguments

data	VPR dataframe from <a href="#">vpr_ctdroi_merge</a> , with calculated variable sigmaT
data_mea	VPR size data frame from <a href="#">vpr_autoid_read</a>
taxa_of_interest	a list of taxa of interest to be included in output dataframe

### Value

A dataframe containing VPR CTD and size data

## Examples

```
data("ctd_roi_merge")
data("roimeas_dat_combine")
category_of_interest = 'Calanus'

ctd_roi_merge$time_hr <- ctd_roi_merge$time_ms /3.6e+06

size_df_f <- vpr_ctdroisize_merge(ctd_roi_merge, data_mea = roimeas_dat_combine,
  taxa_of_interest = category_of_interest)
```

---

vpr_ctdroi_merge	<i>Merge CTD and ROI data from VPR</i>
------------------	--

---

## Description

Combines CTD data (time, hydrographic parameters), with ROI information (identification number) into single dataframe, aligning ROI identification numbers and taxa classifications with time and hydrographic parameters

## Usage

```
vpr_ctdroi_merge(ctd_dat_combine, roi_dat_combine)
```

## Arguments

ctd\_dat\_combine  
a CTD dataframe from VPR processing from [vpr\\_ctd\\_read](#)

roi\_dat\_combine  
a data frame of roi aid data from [vpr\\_autoid\\_read](#)

## Author(s)

E. Chisholm & K. Sorochan

## Examples

```
data('ctd_dat_combine')
data('roi_dat_combine')

ctd_roi_merge <- vpr_ctdroi_merge(ctd_dat_combine, roi_dat_combine)
```

---

vpr\_ctd\_files                      *Create a list of ctd files to be read*

---

**Description**

Searches through typical VP directory structure

**Usage**

```
vpr_ctd_files(castdir, cruise, day_hour)
```

**Arguments**

castdir	root directory for ctd cast files
cruise	cruise name (exactly as in directory structure)
day_hour	vector of day-hour combinations (e.g, dXXX.hXX)

**Details**

Use with caution

**Value**

vector of ctd file paths matching days-hour combinations provided

**Author(s)**

E. Chisholm and K. Sorochan

---

vpr\_ctd\_read                      *Read and format CTD VPR data*

---

**Description**

Acts as a wrapper for [ctd\\_df\\_cols](#)

**Usage**

```
vpr_ctd_read(ctd_files, station_of_interest, day, hour, col_list)
```

**Arguments**

ctd_files	full file paths to vpr ctd .dat files
station_of_interest	VPR station name
day	Day of interest, if not provided will be pulled from file path
hour	Hour of interest, if not provided will be pulled from file path
col_list	Optional list of CTD data column names

**Details**

Reads CTD data and adds day, hour, and station information. Calculates sigma T and depth variables from existing CTD data to supplement raw data. If there are multiple hours of CTD data, combines them into single dataframe.

**WARNING** `ctd_df_cols` is hard coded to accept a specific order of CTD data columns. The names and values in these columns can change based on the specific instrument and should be updated/confirmed before processing data from a new VPR.

**Author(s)**

E. Chisholm & K. Sorochan

**Examples**

```
station_of_interest <- 'test'

ctd_files <- system.file("extdata/COR2019002/rois/vpr5/d222", "h03ctd.dat",
  package = "vpr", mustWork = TRUE)

ctd_dat_combine <- vpr_ctd_read(ctd_files, station_of_interest)
```

---

vpr\_ctd\_ymd *Add Year/ month/ day hour:minute:second information*

---

**Description**

Calculate and record calendar dates for vpr data from day-of-year, hour, and time (in milliseconds) info. Will also add 'time\_hr' parameter if not already present.

**Usage**

```
vpr_ctd_ymd(data, year, offset)
```

**Arguments**

data           VPR data frame from [vpr\\_ctdroi\\_merge](#)  
year           Year of data collection  
offset         time offset in hours between VPR CPU and processed data times (optional)

**Value**

a VPR data frame with complete date/time information in a new row named 'ymdhms'

**Examples**

```
year <- 2019  
data('ctd_roi_merge')  
dat <- vpr_ctd_ymd(ctd_roi_merge, year)
```

---

vpr_day	<i>Get day identifier</i>
---------	---------------------------

---

**Description**

Get day identifier

**Usage**

```
vpr_day(x)
```

**Arguments**

x                 A string specifying the directory and file name of the size file

**Value**

A string of only the day identifier (i.e., "dXXX")

**Author(s)**

K Sorochan

**See Also**

[vpr\\_hour](#), [vpr\\_roi](#), [vpr\\_category](#)

**Examples**

```
day_string <- 'C:/data/cruise/autoid/Calanus/d000/h00'  
vpr_day(day_string)
```



---

vpr_dayhour	<i>Find day &amp; hour info to match each station of interest for processing</i>
-------------	--

---

**Description**

@author E. Chisholm and K. Sorochan

**Usage**

```
vpr_dayhour(stations, file)
```

**Arguments**

stations	a vector of character values naming stations of interest
file	CSV file containing 'day', 'hour', 'station', and 'day_hour' columns

**Value**

Vector of day-hour combinations corresponding to stations of interest

---

vpr_hour	<i>Get hour identifier</i>
----------	----------------------------

---

**Description**

Get hour identifier

**Usage**

```
vpr_hour(x)
```

**Arguments**

x	A string specifying the directory and file name of the size file
---	--

**Value**

A string of only the hour identifier (i.e., "hXX")

**Author(s)**

K Sorochan

**See Also**

[vpr\\_day](#), [vpr\\_roi](#), [vpr\\_category](#)

**Examples**

```
hour_string <- 'C:/data/cruise/autoid/Calanus/d000/h00'
vpr_hour(hour_string)
```

---

vpr\_img\_category      *Explore images by depth and classification*

---

**Description**

Pulls images from specific depth ranges in specific classification group

**Usage**

```
vpr_img_category(
  data,
  min.depth,
  max.depth,
  roiFolder,
  format = "list",
  taxa_of_interest
)
```

**Arguments**

data	data frame containing CTD and ROI data from <a href="#">vpr_ctdroi_merge</a> , which also contains calculated variables sigmaT and time_hr
min.depth	minimum depth of ROIs you are interested in looking at
max.depth	maximum depth of ROIs you are interested in exploring
roiFolder	directory that ROIs are within (can be very general eg. C:/data, but will be quicker to process with more specific file path)
format	option of how images will be output, either as 'list' a list of file names or 'image' where images will be displayed
taxa_of_interest	character string of classification group from which to pull images

---

vpr_img_check	<i>Remove ROI strings from aid and aidmeas files based on a manually organized folder of images</i>
---------------	---

---

**Description**

Should be used after [vpr\\_img\\_copy](#), and manual image removal from created folders

**Usage**

```
vpr_img_check(folder_dir, basepath)
```

**Arguments**

folder_dir	directory path to day hour folders containing manually reorganized images of a specific taxa eg. 'C:/data/cruise_IML2018051/krill/images/' where that folder contains '.....d123.h01/' which contains manually sorted images of krill
basepath	directory path to original Visual Plankton files, specified down to the classification group. eg. 'C:/data/cruise_IML2018051/autoid/krill'

---

vpr_img_copy	<i>Image copying function for specific taxa of interest</i>
--------------	---

---

**Description**

This function can be used to copy images from a particular taxa, day and hour into distinct folders within the auto id directory This is useful for visualizing the ROIs of a particular classification group or for performing manual tertiary checks to remove images not matching classification group descriptions.

**Usage**

```
vpr_img_copy(auto_id_folder, taxas.of.interest, day, hour)
```

**Arguments**

auto_id_folder	eg "D:/VP_data/IML2018051/autoid"
taxas.of.interest	eg. taxas.of.interest <- c('Calanus')
day	character, day of interest
hour	character, hour of interest

---

vpr\_img\_depth      *Explore VPR images by depth bin*

---

### Description

Allows user to pull VPR images from specific depth ranges, to investigate trends before classification of images into taxa groups

### Usage

```
vpr_img_depth(data, min.depth, max.depth, roiFolder, format = "list")
```

### Arguments

data	data frame containing CTD and ROI data from <a href="#">vpr_ctdroi_merge</a> , which also contains calculated variables sigmaT and time_hr
min.depth	minimum depth of ROIs you are interested in looking at
max.depth	maximum depth of ROIs you are interested in exploring
roiFolder	directory that ROIs are within (can be very general eg. C:/data, but will be quicker to process with more specific file path)
format	option of how images will be output, either as 'list' a list of file names or 'image' where images will be displayed

---

vpr\_img\_reclassified      *Explore reclassified images*

---

### Description

Pull image from reclassified or misclassified files produced during [vpr\\_manual\\_classification](#)

### Usage

```
vpr_img_reclassified(day, hour, base_dir, taxa_of_interest, image_dir)
```

### Arguments

day	Character string, 3 digit day of interest of VPR data
hour	Character string, 2 digit hour of interest of VPR data
base_dir	directory path to folder containing day/hour folders in which misclassified and reclassified files are organized (eg. 'C:/VPR_PROJECT/r_project_data_vis/classification files/') which would contain 'd123.h01/reclassified_krill.txt' )
taxa_of_interest	Classification group from which to pull images
image_dir	directory path to ROI images, eg. "E:\\data\\cruise_IML2018051\\", file separator MUST BE "\\" in order to be recognized

**Value**

folders of misclassified or reclassified images inside image\_dir

---

vpr\_manual\_classification

*Function to check results of classification manually*

---

**Description**

Displays each image in day hour specified, prompts user to confirm or deny classification. If classification is denied, asks for a reclassification value based on available taxa

**Usage**

```
vpr_manual_classification(
    day,
    hour,
    basepath,
    taxa_of_interest,
    gr = TRUE,
    scale = "x300",
    opticalSetting = "S2",
    img_bright = TRUE
)
```

**Arguments**

day	day of interest in autoid
hour	hour of interest in autoid
basepath	file path to auto id folder eg 'E:/autoID_EC_07032019/'
taxa_of_interest	list of taxa folders you wish you sort through
gr	logical indicating whether pop up graphic menus are used (user preference - defaults to TRUE)
scale	argument passed to <a href="#">image_scale</a> , default = 'x300'
opticalSetting	specifies optical setting of VPR, defining image frame size, current options are 'S0', 'S1', 'S2' (default), 'S3', see further info in details
img_bright	logical value indicating whether or not to include a blown out high brightness version of image (can be helpful for viewing dark field fine appendages)

**Details**

Optical Setting frame sizes: S0 = 7x7 mm, S1 = 14x14mm, S2 = 24x24mm, S3 = 48x48 mm. These settings define the conversion factor from pixels to millimetres and calculate image size for classification reference

**Development**

- Add "undo" functionality to go back on a typing mistake
- Fix scaling/ size issue so images are consistently sized
- show ROI number for image somewhere for reference when in doubt of classification

---

vpr_oce_create	<i>Create ctd oce object with vpr data</i>
----------------	--

---

**Description**

Formats VPR data frame into oce format CTD object

**Usage**

```
vpr_oce_create(data)
```

**Arguments**

data	data frame of vpr data with variable names 'time_ms', 'fluorescence_mv', 'turbidity_mv', 'n_roi', 'sigmaT'
------	---

**Author(s)**

E. Chisholm

**Examples**

```
data('ctd_roi_merge')  
oce_dat <- vpr_oce_create(ctd_roi_merge)
```

---

vpr_plot_contour	<i>Interpolated contour plot of particular variable</i>
------------------	---

---

**Description**

Creates interpolated contour plot, can be used as a background for ROI or tow yo information

**Usage**

```
vpr_plot_contour(
  data,
  var,
  dup = "mean",
  method = "interp",
  labels = TRUE,
  bw = 1,
  cmo
)
```

**Arguments**

data	data frame needs to include time_hr, depth, and variable of choice (var)
var	variable in dataframe which will be interpolated and plotted
dup	if method == 'interp'. Method of handling duplicates in interpolation, passed to interp function (options: 'mean', 'strip', 'error')
method	Specifies interpolation method, options are 'interp' or 'oce', oce uses slightly different method (oce is least error prone)
labels	logical value indicating whether or not to plot contour labels
bw	bin width defining interval at which contours are labelled
cmo	name of a cmocean plotting theme, see ?cmocean for more information

**Author(s)**

E. Chisholm & Kevin Sorochan

---

vpr\_plot\_histsize      *Plot size frequency histogram*

---

**Description**

Plot size frequency histogram

**Usage**

```
vpr_plot_histsize(data, param, title = NULL, bw = 0.1, xlim = NULL)
```

**Arguments**

data	size data from auto_measure_mm subset into taxa
param	size parameter of interest (corresponds to sub lists within data argument)
title	main title for plot, if left null will default based on parameter and taxa
bw	bin width, defines width of bars on histogram, defaults to 0.1, decrease for more detail
xlim	plot xlim, defaults to min max of data if not provided

**Note**

param options are typically 'Perimeter', 'Area', 'width1', 'width2', 'width3', 'short\_axis\_length', 'long\_axis\_length'

**Author(s)**

E. Chisholm

---

vpr_plot_profile	<i>Plots VPR profiles of temperature, salinity, density, fluorescence and concentration (by classification group)</i>
------------------	---

---

**Description**

This plot allows a good overview of vertical distribution of individual classification groups along with reference to hydrographic parameters. Facet wrap is used to create distinct panels for each taxa provided

**Usage**

```
vpr_plot_profile(taxa_conc_n, taxa_to_plot, plot_conc)
```

**Arguments**

taxa_conc_n	A VPR data frame with hydrographic and concentration data separated by taxa (from <a href="#">vpr_roi_concentration</a> )
taxa_to_plot	The specific classification groups which will be plotted, if NULL, will plot all taxa combined
plot_conc	Logical value whether or not to include a concentration plot (FALSE just shows CTD data)

**Value**

A gridded object of at least 3 ggplot objects



---

vpr\_plot\_sizefreq      *Size Frequency plots for VPR data*

---

**Description**

This uses the `hist` plot function in base R to give a histogram of size (long axis length) frequency within a taxa. **!!WARNING:** this function uses hard coded plot attributes

**Usage**

```
vpr_plot_sizefreq(x, number_of_classes, colour_of_bar)
```

**Arguments**

`x`                      a data frame with columns 'taxa', 'long\_axis\_length'  
`number_of_classes`      numeric value passed to `nclass` argument in `hist()`  
`colour_of_bar`          character value defining colour of plotted bars

**Author(s)**

K. Sorochan

---

vpr\_plot\_TS              *Make a balloon plot against a TS plot*

---

**Description**

TS balloon plot with ROI concentration, sorted by taxa includes isopycnal line calculations

**Usage**

```
vpr_plot_TS(x, reference.p = 0, var)
```

**Arguments**

`x`                      dataframe with temperature, salinity, number of rois (`n_roi_bin`)  
`reference.p`          reference pressure (default at 0 for surface)- used to calculate isopycnals  
`var`                    variable on which size of points will be based, eg `conc_m3` or `n_roi_bin`

**Note**

modified from source: [https://github.com/Davidatlarge/ggTS/blob/master/ggTS\\_DK.R](https://github.com/Davidatlarge/ggTS/blob/master/ggTS_DK.R)

**Author(s)**

E. Chisholm

---

vpr\_plot\_TScat            *Make a balloon plot*

---

### Description

Balloon plot against a TS plot with ROI concentration and sorted by taxa includes isopycnal line calculations. Version of [vpr\\_plot\\_TS](#), with only relevant\* taxa specified. \*to current analysis and research objectives (See note).

### Usage

```
vpr_plot_TScat(x, reference.p = 0)
```

### Arguments

x                        dataframe with temperature, salinity, number of rois named by taxa  
reference.p            reference pressure (default at 0 for surface)- used to calculate isopycnals

### Note

**WARNING HARD CODED FOR 5 TAXA, CALANUS, KRILL, ECHINODERM LARVAE, SMALL COPEPOD, CHAETOGNATHS !!** Uses isopycnal labelling method which does not label every contour

modified from source: [https://github.com/Davidatlarge/ggTS/blob/master/ggTS\\_DK.R](https://github.com/Davidatlarge/ggTS/blob/master/ggTS_DK.R)

---

vpr\_pred\_read            *Read prediction output from a CNN model*

---

### Description

Read prediction output from a CNN model

### Usage

```
vpr_pred_read(filename)
```

### Arguments

filename                model prediction output file (.txt) from vpr\_transferlearn::save\_output()

### Value

a dataframe

---

vpr_roi	<i>Get roi ids from string</i>
---------	--------------------------------

---

**Description**

Get roi ids from string

**Usage**

```
vpr_roi(x)
```

**Arguments**

x                    A string specifying directory and file name of roi

**Value**

A string of only the 10 digit roi identifier

**Author(s)**

K Sorochan

**See Also**

[vpr\\_hour](#), [vpr\\_day](#), [vpr\\_category](#)

**Examples**

```
roi_string <- 'roi.0100000000.tif'  
vpr_roi(roi_string)
```

---

vpr_roi_concentration	<i>Calculate VPR concentrations</i>
-----------------------	-------------------------------------

---

**Description**

Calculates concentrations for each named taxa in dataframe

**Usage**

```
vpr_roi_concentration(
  data,
  taxas_list,
  station_of_interest,
  binSize,
  imageVolume
)
```

**Arguments**

data	a VPR dataframe as produced by <a href="#">vpr_ctdroi_merge</a>
taxas_list	a list of character strings representing taxa present in the station being processed
station_of_interest	The station being processed
binSize	passed to <a href="#">bin_calculate</a> , determines size of depth bins over which data is averaged
imageVolume	the volume of VPR images used for calculating concentrations (mm <sup>3</sup> )

**Examples**

```
data('ctd_roi_merge')
ctd_roi_merge$time_hr <- ctd_roi_merge$time_ms /3.6e+06

taxas_list <- c('Calanus', 'krill')
binSize <- 5
station_of_interest <- 'test'
imageVolume <- 83663

taxa_conc_n <- vpr_roi_concentration(ctd_roi_merge, taxas_list,
station_of_interest, binSize, imageVolume)
```

---

vpr\_save

*Save VPR data as an [as.oce](#) object*


---

**Description**

Save VPR data as an [as.oce](#) object

**Usage**

```
vpr_save(data, metadata)
```

**Arguments**

data	a VPR data frame
metadata	(optional) a named list of character values giving metadata values. If this argument is not provided user will be prompted for a few generic metadata requirements.

**Details**

This function will pass a VPR data frame object to an oce object. Using an oce object as the default export format for VPR data allows for metadata and data to be kept in the same, space efficient file, and avoid redundancy in the data frame. The function check for data parameters that may actually be metadata parameters (rows which have the same value repeated for every observation). These parameters will automatically be copied into the metadata slot of the oce object. The function will also prompt for a variety of required metadata fields. Depending on specific research / archiving requirements, these metadata parameters could be updated by providing the argument metadata.

Default metadata parameters include 'deploymentType', 'waterDepth', 'serialNumber', 'latitude', 'longitude', 'castDate', 'castStartTime', 'castEndTime', 'processedBy', 'opticalSetting', 'imageVolume', 'comment'.

**Value**

an oce CTD object with all VPR data as well as metadata

**Examples**

```
data("taxa_conc_n")
metadata <- c('deploymentType' = 'towyo', 'waterDepth' =
max(ctd_roi_merge$pressure), 'serialNumber' = NA, 'latitude' = 47,
'longitude' = -65, 'castDate' = '2019-08-11', 'castStartTime' = '00:00',
'castEndTime' = '01:00', 'processedBy' = 'E. Chisholm', 'opticalSetting' =
'S2', 'imageVolume' = 83663, 'comment' = 'test data')

oce_dat <- vpr_save(taxa_conc_n, metadata)
# save(oce_dat, file = vpr_save.RData') # save data
```

---

vpr\_size\_bin

*Bin VPR size data*


---

**Description**

Calculates statistics for VPR measurement data in depth averaged bins for analysis and visualization

**Usage**

```
vpr_size_bin(data_all, bin_mea)
```

**Arguments**

data_all	a VPR CTD and measurement dataframe from <a href="#">vpr_ctdroisize_merge</a>
bin_mea	Numerical value representing size of depth bins over which data will be combined, unit is metres, typical values range from 1 - 5

**Value**

a dataframe of binned VPR size data statistics including number of observations, median, interquartile ranges, salinity and pressure, useful for making boxplots

**Examples**

```
data('size_df_f')
vpr_size_bin(size_df_f, bin_mea = 5)
```

---

vpr\_summary

*Data Summary Report*


---

**Description**

Part of VP easy plot processing, prints data summary report to give quantitative, exploratory analysis of data

**Usage**

```
vpr_summary(all_dat, fn, tow = tow, day = day, hour = hour)
```

**Arguments**

all_dat	data frame containing VPR and CTD data including time_ms, time_hr, conductivity, temperature, pressure, salinity, fluorescence_mv, turbidity_mv, sigmaT
fn	file name to save data summary, if not provided, summary will print to console
tow	VPR tow number
day	julian day
hour	two digit hour (24 hr clock)

**Author(s)**

E Chisholm

---

vpr_trrois_size	<i>Get size data from idsize files</i>
-----------------	--

---

**Description**

useful for getting size distribution of known rois from each taxa. gathers size information from idsize text files produced when training a new classifier in VP (Visual Plankton)

**Usage**

```
vpr_trrois_size(directory, taxa, opticalSetting)
```

**Arguments**

directory	cruise directory eg. 'C:/data/IML2018051/'
taxa	list of character elements containing taxa of interest
opticalSetting	VPR optical setting determining conversion between pixels and millimetres (options are 'S0', 'S1', 'S2', or 'S3')

---

vp_plot_matrix	<i>Plots normalized confusion matrix</i>
----------------	--

---

**Description**

Plots normalized confusion matrix

**Usage**

```
vp_plot_matrix(cm, classes, type, addLabels = TRUE, threshold = NULL)
```

**Arguments**

cm	Confusion matrix (numeric)
classes	character list of classes present in confusion matrix (ordered)
type	character value 'NN', 'SVM' or 'Dual', appended to 'Confusion Matrix' to create title
addLabels	logical value whether to add percentage accuracy labels to plot (defaults to TRUE)
threshold	numeric value which determines the minimum value of frequency labelled on the plot on a normalized scale of 0-1 (useful for highlighting significant disagreement)

**Value**

a visualization of the confusion matrix, normalized

**Author(s)**

E. Chisholm

---

vp_plot_unkn	<i>Function to visualize losses to unknown category due to disagreement in Dual classifier</i>
--------------	--

---

**Description**

Makes confusion matrix like plot, where x axis represent SVM classification, y axis represent NN classification Allows visual summary of data lost to unknown category

**Usage**

```
vp_plot_unkn(cm, classes, threshold = 0, summary = TRUE, sample_size = NULL)
```

**Arguments**

cm	dual unknown confusion matrix from VP
classes	taxa groups in order, from VP
threshold	minimum value which will be labelled in plot
summary	logical to add text summary to plot E. Chisholm May 2019
sample_size	character string describes the sample size used to train the model being plotted (optional)



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