

# pastclim: : CHEAT SHEET



## Basics

*pastclim* is an **R package** to easily **download and manipulate paleoclimatic data**.

### INSTALLATION

Installation through devtools

```
library(devtools)
install_github("EvolEcolGroup/pastclim",
  build_vignettes = TRUE)
```

### VIGNETTE

```
vignette("pastclim_overview",
  package = "pastclim")
```

## Data

### NATIVE DATASETS

#### Beyer2020:

- 22 variables
- present-120k years ago
- every 1000/2000 years

#### Krapp2021:

- 22 variables
- present-800k years ago
- every 1000 years

#### Example:

- 3 variables
- 3 time slices

Do not use for analyses!

### LIST ALL AVAILABLE DATASETS

```
vignette("available_datasets",
  package = "pastclim")
```

### ADD MORE DATA

Follow the instructions from the **“custom datasets” vignette**

```
vignette("custom_datasets",
  package = "pastclim")
```

## Download the data

Download the full dataset

By default it will be stored within the package



```
download_dataset(dataset="Beyer2020")
```

Download only a few variables



```
download_dataset(dataset="Beyer2020",
  bio_variables = c("bio01", "bio05"))
```

Download a dataset into a path of choice



```
download_dataset(dataset="Beyer2020",
  path_to_nc = "~/paleoclimate")
```

## Working with locations

Retrieve the climate for several locations scattered in time and/or space.

Format them as a data.frame with longitude, latitude, age.



```
locations <- data.frame(
  x=c(0,90,-120,-7),      # longitude
  y=c(20,45,60,37),      # latitude
  time=c(0,-10000, -20000, -10000))
```

Retrieve the climate



```
location_slice(
  x= locations[,c("x", "y")],
  time_bp = locations$time,
  bio_variables=c("bio01", "bio12"),
  dataset="Example")
```

**WARNING:** Some locations may fall under ice or water, where the climate is unavailable. `nn_interpol = TRUE` (the default) interpolates the climate among the neighbouring cells.

Retrieve time series of the climate for each location (interpolation not available)



```
location_series(
  x= locations[,c("x", "y")],
  bio_variables=c("bio01", "bio12"),
  dataset="Example")
```

## Working with regions

Get climate for the world



```
region_slice(
  time_bp = -20000,
  bio_variables=c("bio01", "bio12"),
  dataset = "Example")
```

Get time series for the world



```
region_series(
  bio_variables=c("bio01", "bio12"),
  dataset = "Example")
```

Both functions can include the arguments

- ‘crop’ to crop results based on a polygon
- ‘ext’ to crop results based on an extent

## Cropping

Eight preset extents are available in *pastclim* and can be listed through the command



```
names(region_extent)
```

- "Africa" • "America" • "Asia"
- "Europe" • "Eurasia" • "N\_America"
- "S\_America" • "Oceania"

Get time series for Asia using an extent



```
region_series(
  bio_variables=c("bio01", "bio12"),
  dataset = "Example",
  ext=region_extent$Asia)
```

*pastclim* includes five pre-generated masks for the main continental masses



```
names(region_outline)
```

- "Africa" • "North\_America" • "Eurasia"
- "Oceania" • "South\_America"

Get climate from Africa using a mask



```
region_slice(
  time_bp = -20000,
  c("bio01", "bio12"),
  dataset = "Example",
  crop = region_outline$Africa)
```

## Sampling

Sampling 100 random points from a region

```
climate_20k <- region_slice(
  time_bp = -20000,
  bio_variables = c("bio01", "bio10"),
  dataset = "Example")
```

```
sample_region_slice(
  climate_20k,
  size = 100)
```

Sampling from a time series, 100 points from the first time slice, 50 from the second.

```
climate_ts <- region_series(
  time_bp = c(-20000, -10000),
  bio_variables = c("bio10", "bio12"),
  dataset = "Example")
```

```
sample_region_series(
  climate_ts,
  size = c(100, 50))
```

To sample the same amount from all time slices write only one number

## Citation

Cite *pastclim*

```
citation("pastclim")
```

Cite database

```
help("Beyer2020")
```

## Need help?



<https://rdrr.io/github/EvolEcolGroup/pastclim/man/>



<https://github.com/EvolEcolGroup/pastclim>



<https://evolecolgroup.github.io/pastclim/>



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