



Analyzing space-time satellite data for disease ecology applications with **GRASS GIS** and R stats

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Applications of RS & GIS for disease ecology

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Disease Ecology, Rodents, Hantavirus

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GRASS and for disease ecology



rgrass7

- `initGRASS()`: starts a GRASS GIS session from R
- `execGRASS()`: executes GRASS GIS commands
- `gmeta()`: shows GRASS location metadata
- `readVECT()` and `readRAST()`: read vector and raster maps from GRASS into `sf/sp` objects
- `writeVECT()` and `writeRAST()`: write `sf/sp` objects into GRASS GIS database

An update to support stars for GRASS raster maps is on the way! Kudos to Roger Bivand!



GRASS GIS and can be used together in two ways:

- Using R within a GRASS GIS session
- Using GRASS GIS within an R session

Details and examples at the [GRASS and R wiki](#)

- Using **R within a GRASS GIS session**, i.e. starting R (or RStudio) from the GRASS GIS terminal
 - we do not need to initialize GRASS with `initGRASS()`, just type `R` or `rstudio` & in the GRASS GIS terminal
 - we access GRASS GIS functionalities and database through `execGRASS()`
 - we use `readVECT()`, `readRAST()` to read data from GRASS DB to do analysis and plots in R
 - we write data (back) to GRASS GIS database with `writeVECT()` and `writeRAST()`



- Using **GRASS GIS within an R session**, i.e. we connect to GRASS GIS database from within R (or RStudio).
 - we need to start GRASS GIS with `initGRASS()` (usually by creating a throw away location)
 - we access GRASS GIS functionalities through `execGRASS()` (usually to apply them on data outside GRASS DB)



For more detailed examples check:

- R and GRASS presentation
- Example of GRASS - R for raster time series wiki



link2GI

See the [vignette](#) on how to set GRASS database with link2GI
for further details



Hands-on to space-time analysis for disease ecology with GRASS

GIS and R



Overview

- Import species records
- Create random background points
- Create environmental layers
- Read data into R
- Model species distribution
- Model evaluation
- Visualization of results

Check [Analyzing space-time satellite data with GRASS GIS for environmental monitoring](#) presentation for an intro to time series in GRASS GIS.

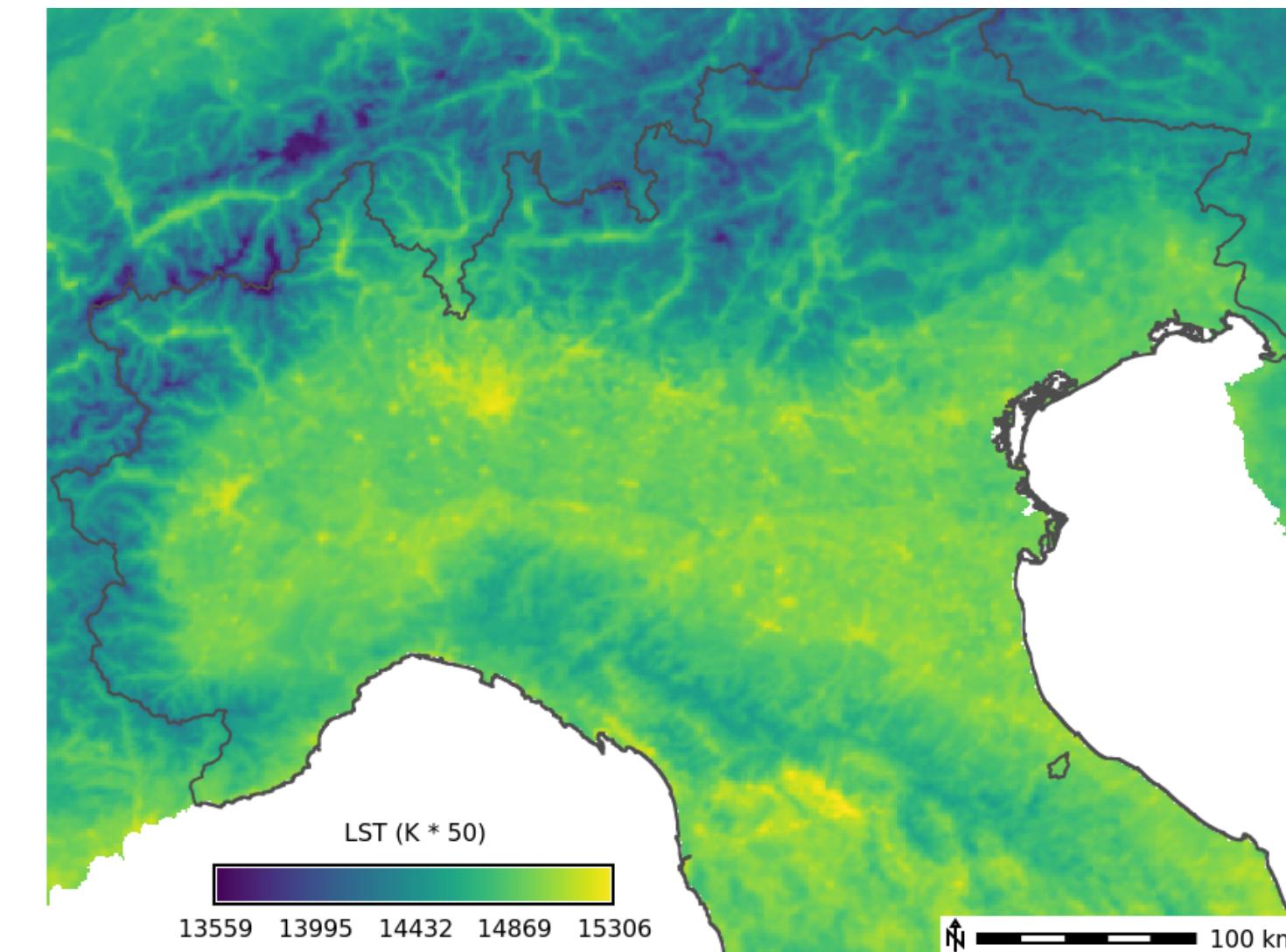
Data for the session

- Records of *Aedes albopictus* (Asian tiger mosquito) in Northern Italy from [GBIF](#)
- Environmental layers derived from reconstructed daily MODIS LST products



Data for the session: LST

- MODIS LST
reconstructed by
mundialis based on
Metz et al. 2017
- Daily average LST
- 1 km spatial
resolution
- Converted to Celsius
degrees



LST, July 2018 - Northern Italy.

 get sample location and code 

- eu_laea location with LST mapset: download and unzip within your grassdata folder
- GRASS script
- R script



Let's start GRASS GIS!



Importing species records and creating random background points

```
#!/bin/bash
#####
# GRASS GIS commands for the session:
# "Analysis of space-time satellite data for disease ecology
# applications with GRASS GIS and R stats" at OpenGeoHub Summer School,
# Muenster (Germany)
#
# Modelling Aedes albopictus potential distribution in Northern Italy
#
# Author: Veronica Andreo
#
# Date: August, 2019
#####
#
# Download data from GBIF for Aedes albopictus in Northern Italy
#
# Set computational region
```

Importing species records and creating random background points

```
# Date: August, 2019
#####
#
# Download data from GBIF for Aedes albopictus in Northern Italy
#
#
# Set computational region
g.region -p raster=lst_2014.001_avg

# Install extension (requires pygbif: pip install pygbif)
g.extension extension=v.in.pygbif

# Import data from GBIF
v.in.pygbif output=aedes_albopictus \
taxa="Aedes albopictus" \
date_from="2014-01-01" \
date_to="2018-12-31"
```

Importing species records and creating random background points

```
# # Download data from GBIF for Aedes albopictus in Northern Italy
#
# Set computational region
g.region -p raster=lst_2014.001_avg

# Install extension (requires pygbif: pip install pygbif)
g.extension extension=v.in.pygbif

# Import data from GBIF
v.in.pygbif output=aedes_albopictus \
  taxa="Aedes albopictus" \
  date_from="2014-01-01" \
  date_to="2018-12-31"
```

Importing species records and creating random background points

```
# Set computational region
g.region -p raster=lst_2014.001_avg

# Install extension (requires pygbif: pip install pygbif)
g.extension extension=v.in.pygbif

# Import data from GBIF
v.in.pygbif output=aedes_albopictus \
  taxa="Aedes albopictus" \
  date_from="2014-01-01" \
  date_to="2018-12-31"

#
# Create background points
#
# Create buffer around Aedes albopictus records
```

Importing species records and creating random background points

```
date_from="2014-01-01" \
date_to="2018-12-31"

#
# Create background points
#


# Create buffer around Aedes albopictus records
v.buffer input=aedes_albopictus \
          output=aedes_buffer \
          distance=1000

# Create a vector mask to limit background points
r.mapcalc \
  expression="rast_mask = if(lst_2014.001_avg, 1, null())"

r.to.vect input=rast_mask \
          output=vect_mask \
          type=area
```

Importing species records and creating random background points

```
# Create buffer around Aedes albopictus records
v.buffer input=aedes_albopictus \
    output=aedes_buffer \
    distance=1000

# Create a vector mask to limit background points
r.mapcalc \
    expression="rast_mask = if(lst_2014.001_avg, 1, null())"

r.to.vect input=rast_mask \
    output=vect_mask \
    type=area

# Subtract buffers from vector mask
v.overlay ainput=vect_mask \
    binput=aedes_buffer \
    operator=xor \
    output=mask_background
```

Importing species records and creating random background points

```
# Create a vector mask to limit background points
r.mapcalc \
    expression="rast_mask = if(lst_2014.001_avg, 1, null())"

r.to.vect input=rast_mask \
    output=vect_mask \
    type=area

# Subtract buffers from vector mask
v.overlay ainput=vect_mask \
    binput=aedes_buffer \
    operator=xor \
    output=mask_background

# Generate random background points
v.random output=background_points \
    npoints=1000 \
    restrict=mask_background \
    seed=3749
```

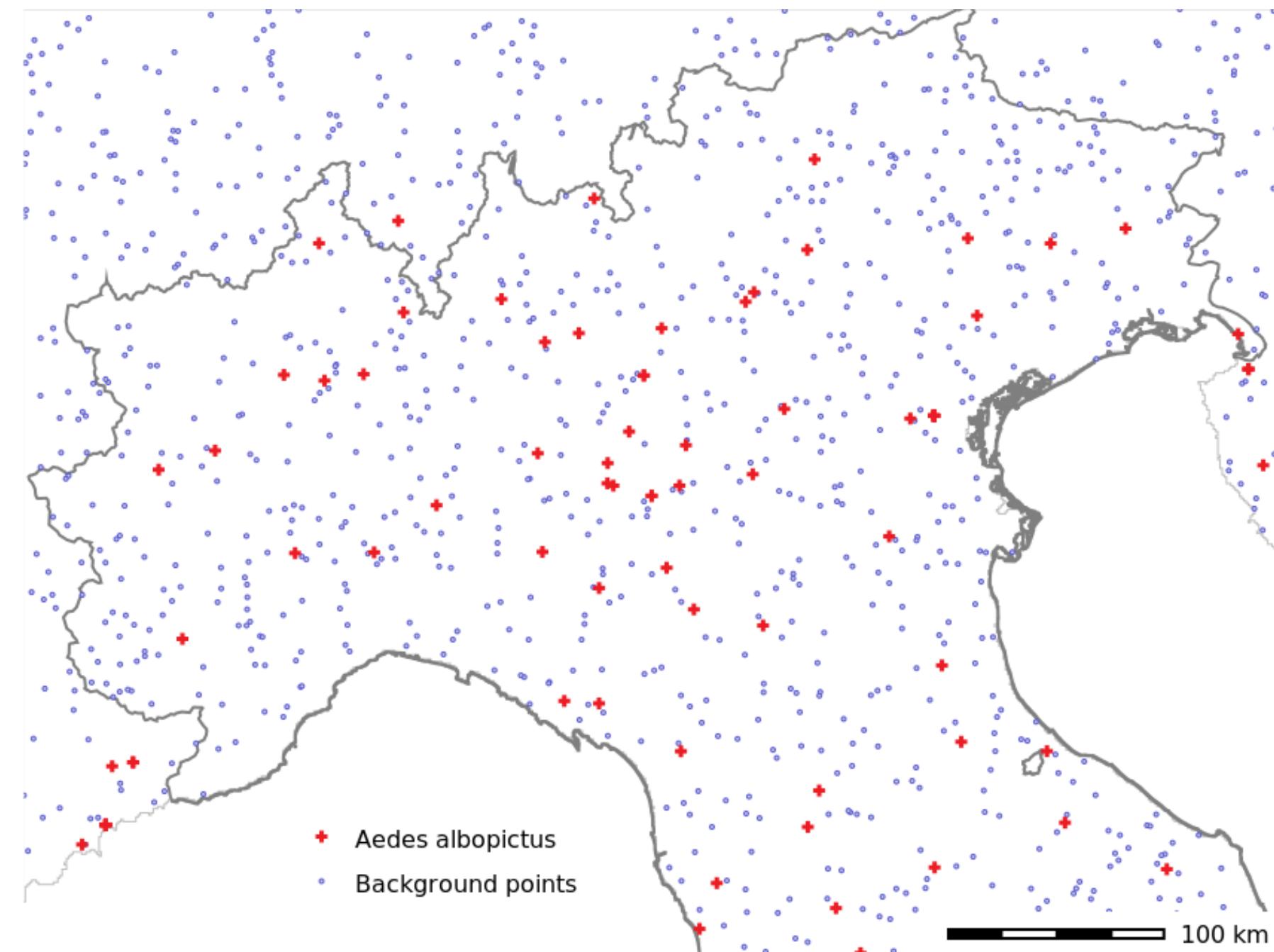
Importing species records and creating random background points

```
type=area

# Subtract buffers from vector mask
v.overlay ainput=vect_mask \
    binput=aedes_buffer \
    operator=xor \
    output=mask_background

# Generate random background points
v.random output=background_points \
    npoints=1000 \
    restrict=mask_background \
    seed=3749

#
# Create daily LST STRDS
#
```



Create daily LST STRDS

```
#!/bin/bash
#####
# GRASS GIS commands for the session:
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#
# Download data from GBIF for Aedes albopictus in Northern Italy
#
#
# Set computational region
```

Create daily LST STRDS

```
restrict=mask_background \
seed=3749

#
# Create daily LST STRDS
# 

# Create time series
t.create type=strds \
temporaltype=absolute \
output=lst_daily \
title="Average Daily LST" \
description="Average daily LST in degree C - 2014-2018"

# Get list of maps
g.list type=raster \
pattern="lst_201*" \
output=list_lst.csv
```

Create daily LST STRDS

```
# Create time series  
t.create type=strds \  
temporaltype=absolute \  
output=lst_daily \  
title="Average Daily LST" \  
description="Average daily LST in degree C - 2014-2018"  
  
# Get list of maps  
g.list type=raster \  
pattern="lst_201*" \  
output=list_lst.csv  
  
# Register maps in strds  
t.register -i input=lst_daily \  
file=list_lst.csv \  
increment="1 days" \  
start="2014-01-01"
```

Create daily LST STRDS

```
temporalitytype=absolute \
output=lst_daily \
title="Average Daily LST" \
description="Average daily LST in degree C - 2014-2018"

# Get list of maps
g.list type=raster \
pattern="lst_201*" \
output=list_lst.csv

# Register maps in strds
t.register -i input=lst_daily \
file=list_lst.csv \
increment="1 days" \
start="2014-01-01"

# Get info about the strds
t.info input=lst_daily

#
```

Create daily LST STRDS

```
# Get list of maps
g.list type=raster \
    pattern="lst_201*" \
    output=list_lst.csv

# Register maps in strds
t.register -i input=lst_daily \
    file=list_lst.csv \
    increment="1 days" \
    start="2014-01-01"

# Get info about the strds
t.info input=lst_daily

#
# Generate environmental variables from LST STRDS
#
```

Generate environmental variables from LST STRDS

```
#!/bin/bash
#####
# GRASS GIS commands for the session:
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# applications with GRASS GIS and R stats" at OpenGeoHub Summer School,
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#####

#
# Download data from GBIF for Aedes albopictus in Northern Italy
#
#
# Set computational region
```

Generate environmental variables from LST STRDS

```
# Long term monthly avg, min and max LST
for i in $(seq -w 1 12) ; do

    # average
    t.rast.series input=lst_daily \
        method=average \
        where="strftime('%m', start_time)='${i}'" \
        output=lst_average_${i}

    # minimum
    t.rast.series input=lst_daily \
        method=minimum \
        where="strftime('%m', start_time)='${i}'" \
        output=lst_minimum_${i}

    # maximum
    t.rast.series input=lst_daily \
        method=maximum \
        where="strftime('%m', start_time)='${i}'" \
        output=lst_maximum_${i}
```

Generate environmental variables from LST STRDS

Generate environmental variables from LST STRDS

```
t.rast.series input=lst_daily \
  method=maximum \
  where="strftime('%m', start_time)=' ${i}'" \
  output=lst_maximum_${i}

done

# Install extension
g.extension extension=r.bioclim

# Estimate temperature related bioclimatic variables

r.bioclim \
  tmin=$(g.list type=raster pattern="lst_minimum_??" separator=",") \
  tmax=$(g.list type=raster pattern="lst_maximum_??" separator=",") \
  tavg=$(g.list type=raster pattern="lst_average_??" separator=",") \
  output=worldclim_

## Spring warming
```

Generate environmental variables from LST STRDS

```
r.bioclim \
tmin=$(g.list type=raster pattern="lst_minimum_??" separator=",") \
tmax=$(g.list type=raster pattern="lst_maximum_??" separator=",") \
tavg=$(g.list type=raster pattern="lst_average_??" separator=",") \
output=worldclim_ \
## Spring warming
# Annual spring warming: slope(daily Tmean february-march-april)
t.rast.aggregate input=lst_daily \
output=annual_spring_warming \
basename=spring_warming \
suffix=gran \
method=slope \
granularity="1 years" \
where="strftime('%m',start_time)='02' or \
strftime('%m',start_time)='03' or \
strftime('%m', start_time)='04'"
```

Generate environmental variables from LST STRDS

```
# Annual spring warming: slope(daily Tmean february–march–april)
t.rast.aggregate input=lst_daily \
    output=annual_spring_warming \
    basename=spring_warming \
    suffix=gran \
    method=slope \
    granularity="1 years" \
    where="strftime('%m',start_time)='02' or \
        strftime('%m',start_time)='03' or \
        strftime('%m', start_time)='04'"'

# Average spring warming
t.rast.series input=annual_spring_warming \
    output=avg_spring_warming \
    method=average

## Autumnal cooling

# Annual autumnal cooling: slope(daily Tmean august–september–october)
```

Generate environmental variables from LST STRDS

```
strftime('%m', start_time)='04'"

# Average spring warming
t.rast.series input=annual_spring_warming \
    output=avg_spring_warming \
    method=average

## Autumnal cooling

# Annual autumnal cooling: slope(daily Tmean august-september-october)
t.rast.aggregate input=lst_daily \
    output=annual_autumnal_cooling \
    basename=autumnal_cooling \
    suffix=gran \
    method=slope \
    granularity="1 years" \
    where="strftime('%m',start_time)='08' or \
        strftime('%m',start_time)='09' or \
        strftime('%m', start_time)='10'"
```

Generate environmental variables from LST STRDS

```
# Annual autumnal cooling: slope(daily Tmean august-september-october)
t.rast.aggregate input=lst_daily \
    output=annual_autumnal_cooling \
    basename=autumnal_cooling \
    suffix=gran \
    method=slope \
    granularity="1 years" \
    where="strftime('%m',start_time)='08' or \
        strftime('%m',start_time)='09' or \
        strftime('%m', start_time)='10'"

# Average autumnal cooling
t.rast.series input=annual_autumnal_cooling \
    output=avg_autumnal_cooling \
    method=average

## Start/end and length of mosquito growing season
```

Generate environmental variables from LST STRDS

Generate environmental variables from LST STRDS

```
Average annual cooling  
t.rast.series input=annual_autumnal_cooling \  
output=avg_autumnal_cooling \  
method=average  
  
## Start/end and length of mosquito growing season  
  
# Install extension  
g.extension extension=r.seasons  
  
# Detect seasons  
for YEAR in $(seq 2014 2018) ; do  
  
    # Get map list per year  
    t.rast.list -u lst_daily \  
    columns=name \  
    where="strftime('%Y',start_time)='${YEAR}'" \  
    output=list_${YEAR}.txt  
  
    # Mosquito season (threshold: 10C, min duration: 150 days)  
    r.seasons file=list_${YEAR}.txt \  
    --threshold 10 --min-duration 150
```

Generate environmental variables from LST STRDS

```
g.extension extension=seasons

# Detect seasons
for YEAR in $(seq 2014 2018) ; do

    # Get map list per year
    t.rast.list -u lst_daily \
        columns=name \
        where="strftime('%Y',start_time)='${YEAR}'" \
        output=list_${YEAR}.txt

    # Mosquito season (threshold: 10C, min duration: 150 days)
    r.seasons file=list_${YEAR}.txt \
        prefix=mosq_season_${YEAR}_ \
        n=1 \
        nout=mosq_season_${YEAR} \
        threshold_value=10 \
        min_length=150 \
        max_gap=12

    # Length core season
```

Generate environmental variables from LST STRDS

```
# Mosquito season (threshold: 10C, min duration: 150 days)
r.seasons file=list_${YEAR}.txt \
prefix=mosq_season_${YEAR}_ \
n=1 \
nout=mosq_season_${YEAR} \
threshold_value=10 \
min_length=150 \
max_gap=12

# Length core season
r.mapcalc expression="mosq_season_length_${YEAR}_1 = \
mosq_season_${YEAR}_1_end1 - mosq_season_${YEAR}_1_start1 + 1"

# Length extended season
r.mapcalc expression="mosq_season_length_${YEAR}_2 = \
mosq_season_${YEAR}_1_end2 - mosq_season_${YEAR}_1_start2 + 1"

done
```

Generate environmental variables from LST STRDS

```
# Length core season
r.mapcalc expression="mosq_season_length_${YEAR}_1 = \
mosq_season_${YEAR}_1_end1 - mosq_season_${YEAR}_1_start1 + 1"

# Length extended season
r.mapcalc expression="mosq_season_length_${YEAR}_2 = \
mosq_season_${YEAR}_1_end2 - mosq_season_${YEAR}_1_start2 + 1"

done

# Average length of mosquito season
r.series \
  input=$(g.list type=raster pattern=mosq_season_length*1 separator=",") \
  output=avg_mosq_season_1_length \
  method=average

r.series \
  input=$(g.list type=raster pattern=mosq_season_length*2 separator=",") \
  output=avg_mosq_season_2_length \
  method=average
```

Generate environmental variables from LST STRDS

```
r.series \
    input=$(g.list type=raster pattern=mosq_season_length*1 separator=",") \
    output=avg_mosq_season_1_length \
    method=average

r.series \
    input=$(g.list type=raster pattern=mosq_season_length*2 separator=",") \
    output=avg_mosq_season_2_length \
    method=average

## Number of days with LSTmean >= 20 and <= 30

# Keep only pixels meeting the condition
t.rast.algebra -n \
    expression="tmean_higher20_lower30 = if(lst_daily >= 20.0 && \
lst_daily <= 30.0, 1, null())" \
    basename=tmean_higher20_lower30 \
    suffix=gran
```

Generate environmental variables from LST STRDS

```
method=average

## Number of days with LSTmean >= 20 and <= 30

# Keep only pixels meeting the condition
t.rast.algebra -n \
  expression="tmean_higher20_lower30 = if(lst_daily >= 20.0 && \
lst_daily <= 30.0, 1, null())" \
  basename=tmean_higher20_lower30 \
  suffix=gran

# Count how many times per year the condition is met
t.rast.aggregate input=tmean_higher20_lower30 \
  output=count_tmean_higher20_lower30 \
  basename=tmean_higher20_lower30 \
  suffix=gran \
  method=count \
  granularity="1 years"
```

Generate environmental variables from LST STRDS

```
expression="tmean_higher20_lower30 = if(lst_daily >= 20.0 && \
lst_daily <= 30.0, 1, null())" \
basename=tmean_higher20_lower30 \
suffix=gran

# Count how many times per year the condition is met
t.rast.aggregate input=tmean_higher20_lower30 \
output=count_tmean_higher20_lower30 \
basename=tmean_higher20_lower30 \
suffix=gran \
method=count \
granularity="1 years"

# Inspect values
t.rast.list \
count_tmean_higher20_lower30 \
columns=name,start_time,end_time,min,max

# Average number of days with LSTmean >= 20 and <= 30
t.rast.series input=count_tmean_higher20_lower30 \
```

Generate environmental variables from LST STRDS

```
# Count how many times per year the condition is met
t.rast.aggregate input=tmean_higher20_lower30 \
  output=count_tmean_higher20_lower30 \
  basename=tmean_higher20_lower30 \
  suffix=gran \
  method=count \
  granularity="1 years"

# Inspect values
t.rast.list \
  count_tmean_higher20_lower30 \
  columns=name,start_time,end_time,min,max

# Average number of days with LSTmean >= 20 and <= 30
t.rast.series input=count_tmean_higher20_lower30 \
  output=avg_count_tmean_higher20_lower30 \
  method=average

## Number of consecutive days with LSTmean >= 33
```

Generate environmental variables from LST STRDS

```
# Inspect values
t.rast.list \
  count_tmean_higher20_lower30 \
  columns=name,start_time,end_time,min,max

# Average number of days with LSTmean >= 20 and <= 30
t.rast.series input=count_tmean_higher20_lower30 \
  output=avg_count_tmean_higher20_lower30 \
  method=average

## Number of consecutive days with LSTmean >= 33

# Create annual mask
t.rast.aggregate input=lst_daily \
  output=annual_mask \
  basename=annual_mask \
  suffix=gran \
  granularity="1 year" \
  method=count
```

Generate environmental variables from LST STRDS

```
method=average

## Number of consecutive days with LSTmean >= 33

# Create annual mask
t.rast.aggregate input=lst_daily \
    output=annual_mask \
    basename=annual_mask \
    suffix=gran \
    granularity="1 year" \
    method=count

# Replace values by zero
t.rast.mapcalc \
    input=annual_mask \
    output=annual_mask_0 \
    expression="if(annual_mask, 0)" \
    basename=annual_mask_0
```

Generate environmental variables from LST STRDS

```
output=annual_mask \
basename=annual_mask \
suffix=gran \
granularity="1 year" \
method=count

# Replace values by zero
t.rast.mapcalc \
input=annual_mask \
output=annual_mask_0 \
expression="if(annual_mask, 0)" \
basename=annual_mask_0

# Calculate consecutive days with LSTmean >= 33
t.rast.algebra \
expression="higher35_consec_days = annual_mask_0 {+,contains,l} \
if(lst_daily >= 33.0 && lst_daily[-1] >= 33.0 || \
lst_daily[1] >= 33.0 && lst_daily >= 33.0, 1, 0)" \
basename=higher35_days \
suffix=gran \
nproc=7
```

Generate environmental variables from LST STRDS

```
Replace values by zero
t.rast.mapcalc \
    input=annual_mask \
    output=annual_mask_0 \
    expression="if(annual_mask, 0)" \
    basename=annual_mask_0

# Calculate consecutive days with LSTmean >= 33
t.rast.algebra \
    expression="higher35_consec_days = annual_mask_0 {+,contains,l} \
    if(lst_daily >= 33.0 && lst_daily[-1] >= 33.0 || \
    lst_daily[1] >= 33.0 && lst_daily >= 33.0, 1, 0)" \
    basename=higher35_days \
    suffix=gran \
    nproc=7

# Inspect values
t.rast.list input=higher35_consec_days \
    columns=name,start_time,end_time,min,max

# Median number of consecutive days with LSTmean >= 33
```

Generate environmental variables from LST STRDS

```
expression="IT(annual_mask, 0)" \
basename=annual_mask_0

# Calculate consecutive days with LSTmean >= 33
t.rast.algebra \
    expression="higher35_consec_days = annual_mask_0 {+,contains,l} \
    if(lst_daily >= 33.0 && lst_daily[-1] >= 33.0 || \
    lst_daily[1] >= 33.0 && lst_daily >= 33.0, 1, 0)" \
    basename=higher35_days \
    suffix=gran \
    nproc=7

# Inspect values
t.rast.list input=higher35_consec_days \
    columns=name,start_time,end_time,min,max

# Median number of consecutive days with LSTmean >= 33
t.rast.series input=higher35_consec_days \
    output=median_higher35_consec_days \
    method=median
```

Generate environmental variables from LST STRDS

```
suffix=gran \
nproc=7

# Inspect values
t.rast.list input=higher35_consec_days \
  columns=name,start_time,end_time,min,max

# Median number of consecutive days with LSTmean >= 33
t.rast.series input=higher35_consec_days \
  output=median_higher35_consec_days \
  method=median

## Growing Degree Days

# Accumulation of degree days
t.rast.accumulate -n input=lst_daily \
  output=mosq_daily_bedd \
  basename=mosq_daily_bedd \
  suffix=gran \
  start="2014-03-01" stop="2018-10-01" \
  -l "mosq_daily" -o "mosq_daily"
```

Generate environmental variables from LST STRDS

```
output=median_nigher35_consec_days \
method=median

## Growing Degree Days

# Accumulation of degree days
t.rast.accumulate -n input=lst_daily \
output=mosq_daily_bedd \
basename=mosq_daily_bedd \
suffix=gran \
start="2014-03-01" stop="2018-10-01" \
cycle="7 months" offset="5 months" \
method=bedd limits=11,30

# Get basic info
t.info input=mosq_daily_bedd

# Inspect values
t.rast.list input=mosq_daily_bedd \
columnnames=starting_time,end_time,min,max
```

Generate environmental variables from LST STRDS

```
basename=mosq_daily_bedd \
suffix=gran \
start="2014-03-01" stop="2018-10-01" \
cycle="7 months" offset="5 months" \
method=bedd limits=11,30

# Get basic info
t.info input=mosq_daily_bedd

# Inspect values
t.rast.list input=mosq_daily_bedd \
columns=name,start_time,end_time,min,max

## Detection of mosquito generations

# Arrays
cycle=($(seq 1 9))
cycle_beg=($(seq 1 300 2700))
cycle_end=($(seq 300 300 2700))
# Length of the array
```

Generate environmental variables from LST STRDS

```
t.rast.list input=mosq_daily_bedd \
    columns=name,start_time,end_time,min,max

## Detection of mosquito generations

# Arrays
cycle=($(seq 1 9))
cycle_beg=($(seq 1 300 2700))
cycle_end=($(seq 300 300 2700))
# Length of the array
count=${#cycle[@]}

for i in $(seq 1 $count) ; do
    echo "cycle: ${cycle[$i-1]} - ${cycle_beg[$i-1]} ${cycle_end[$i-1]}

    # Identify generations
    t.rast.accdetect input=mosq_daily_bedd \
        occurrence=mosq_occurrence_cycle_${cycle[$i-1]} \
        indicator=mosq_indicator_cycle_${cycle[$i-1]} \
        basename=mosq_cycle_${cycle[$i-1]} \
```

Generate environmental variables from LST STRDS

```
# Identify generations
t.rast.accdetect input=mosq_daily_bedd \
occurrence=mosq_occurrence_cycle_${cycle[$i-1]} \
indicator=mosq_indicator_cycle_${cycle[$i-1]} \
basename=mosq_cycle_${cycle[$i-1]} \
start="2014-03-01" stop="2018-10-01" \
cycle="7 months" offset="5 months" \
range=${cycle_beg[$i-1]},${cycle_end[$i-1]}

# Extract areas that have full generations

# Indicator takes values 1, 2 or 3 for start, middle and end of cycle

t.rast.aggregate input=mosq_indicator_cycle_${cycle[$i-1]} \
output=mosq_cycle${cycle[$i-1]}_yearly \
basename=mosq_c${cycle[$i-1]}_yearly \
granularity="1 year" \
method=maximum \
suffix=gran
```

Generate environmental variables from LST STRDS

```
granularity="1 year" \
method=maximum \
suffix=gran

# Keep only complete generations
t.rast.mapcalc input=mosq_cycle${cycle[$i-1]}_yearly \
output=mosq_cycle${cycle[$i-1]}_yearly_clean \
basename=mosq_clean_c${cycle[$i-1]} \
expression="if(mosq_cycle${cycle[$i-1]}_yearly == 3, \
${cycle[$i-1]}, null())"

# Duration of each mosquito generation

# Beginning
t.rast.aggregate input=mosq_occurrence_cycle_${cycle[$i-1]} \
output=mosq_min_day_cycle${cycle[$i-1]} \
basename=occ_min_day_c${cycle[$i-1]} \
method=minimum \
granularity="1 year" \
suffix=gran
```

Generate environmental variables from LST STRDS

```
t.rast.aggregate input=mosq_occurrence_cycle_${cycle[$i-1]} \
    output=mosq_min_day_cycle${cycle[$i-1]} \
    basename=occ_min_day_c${cycle[$i-1]} \
    method=minimum \
    granularity="1 year" \
    suffix=gran

# End
t.rast.aggregate input=mosq_occurrence_cycle_${cycle[$i-1]} \
    output=mosq_max_day_cycle${cycle[$i-1]} \
    basename=occ_max_day_c${cycle[$i-1]} \
    method=maximum \
    granularity="1 year" \
    suffix=gran

# Difference

t.rast.mapcalc \
    input=mosq_min_day_cycle${cycle[$i-1]},mosq_max_day_cycle${cycle[$i-1]} \
    output=mosq_duration_cycle${cycle[$i-1]} \
    basename=mosq_duration_cycle${cycle[$i-1]} \
    expression="mosq_max_day_cycle${cycle[$i-1]} - "
```

Generate environmental variables from LST STRDS

```
method=maximum \
granularity="1 year" \
suffix=gran

# Difference

t.rast.mapcalc \
    input=mosq_min_day_cycle${cycle[$i-1]},mosq_max_day_cycle${cycle[$i-1]} \
    output=mosq_duration_cycle${cycle[$i-1]} \
    basename=mosq_duration_cycle${cycle[$i-1]} \
    expression="mosq_max_day_cycle${cycle[$i-1]} - \
mosq_min_day_cycle${cycle[$i-1]} + 1"

done

# Maximum number of generations per pixel per year
for i in $(seq 1 5) ; do
    g.list type=raster separator="," pattern="mosq_clean_c*_${i}"
    r.series \
        input=$(g.list type=raster pattern="mosq_clean_c*_${i}" separator=",") \
```

Generate environmental variables from LST STRDS

```
input=mosq_min_day_cycle${cycle[$i]},mosq_max_day_cycle${cycle[$i]} \
output=mosq_duration_cycle${cycle[$i-1]} \
basename=mosq_duration_cycle${cycle[$i-1]} \
expression="mosq_max_day_cycle${cycle[$i-1]} - \
mosq_min_day_cycle${cycle[$i-1]} + 1"

done

# Maximum number of generations per pixel per year
for i in $(seq 1 5) ; do
    g.list type=raster separator="," pattern="mosq_clean_c*_${i}"
    r.series \
        input=$(g.list type=raster pattern="mosq_clean_c*_${i}" separator=",") \
        output=mosq_generations_${i} \
        method=maximum
done

# Median number of generations per pixel
r.series \
    input=$(g.list type=raster pattern="mosq_generations_*" separator=",") \
    output=median_mosa_generations \

```

Generate environmental variables from LST STRDS

```
# Maximum number of generations per pixel per year
for i in $(seq 1 5) ; do
    g.list type=raster separator="," pattern="mosq_clean_c*_${i}"
    r.series \
        input=$(g.list type=raster pattern="mosq_clean_c*_${i}" separator=",") \
        output=mosq_generations_${i} \
        method=maximum
done

# Median number of generations per pixel
r.series \
    input=$(g.list type=raster pattern="mosq_generations_*" separator=",") \
    output=median_mosq_generations \
    method=median

# Median duration of generations per year per pixel
for i in $(seq 1 5) ; do
    g.list type=raster separator="," pattern="mosq_duration_cycle*_${i}"
    r.series \
        input=$(g.list type="raster pattern=mosq_duration_cycle*_${i}" separator=",") \
```

Generate environmental variables from LST STRDS

```
method=median
done

# Median number of generations per pixel
r.series \
  input=$(g.list type=raster pattern="mosq_generations_*" separator=",") \
  output=median_mosq_generations \
  method=median

# Median duration of generations per year per pixel
for i in $(seq 1 5) ; do
  g.list type=raster separator="," pattern="mosq_duration_cycle*_${i}"
  r.series \
    input=$(g.list type="raster pattern=mosq_duration_cycle*_${i}" separator=",") \
    output=mosq_generation_median_duration_${i} \
    method=median
done

# Median duration of generations per pixel
r.series \
  input=$(g.list type=raster pattern="mosa_generation_median_duration *" separator=
```

Generate environmental variables from LST STRDS

```
me@me-OptiPlex-5090:~/Documents$ grass -c -r -t -v -i -o -s -w -x -y -z -z  
done  
  
# Median number of generations per pixel  
r.series \  
    input=$(g.list type=raster pattern="mosq_generations_*" separator=",") \  
    output=median_mosq_generations \  
    method=median  
  
# Median duration of generations per year per pixel  
for i in $(seq 1 5) ; do  
    g.list type=raster separator="," pattern="mosq_duration_cycle*_${i}"  
    r.series \  
        input=$(g.list type="raster pattern=mosq_duration_cycle*_${i}" separator=",") \  
        output=mosq_generation_median_duration_${i} \  
        method=median  
done  
  
# Median duration of generations per pixel  
r.series \  
    input=$(g.list type=raster pattern="mosq generation median duration *" separator=
```

Within R...

```
#####
# R commands for the session:
# "Analysis of space-time satellite data for disease ecology
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#
# Modelling Aedes albopictus potential distribution in Northern Italy
#
# Original example contributed by Carol Garzon Lopez
# Adapted by Veronica Andreo
#
# Date: October, 2018
# Updated: August, 2019
#####
#
# Install and load required packages
#
```

Within R...

```
#  
# Install and load required packages  
#  
  
# Install packages  
install.packages("rgrass7")  
install.packages("raster")  
install.packages("sf")  
install.packages("mapview")  
install.packages("biomod2")  
  
# Load libraries  
library(rgrass7)  
library(raster)  
library(sf)  
library(mapview)  
library(biomod2)
```

Within R...

```
# Install packages
install.packages("rgrass7")
install.packages("raster")
install.packages("sf")
install.packages("mapview")

install.packages("biomod2")

# Load libraries
library(rgrass7)
library(raster)
library(sf)
library(mapview)
library(biomod2)

#
# Read vector data
#
```

Within R...

```
library(raster)
library(sf)
library(mapview)
library(biomod2)

#
# Read vector data
#


# Use sf for vectors
use_sf()

# Read vector layers
Aa_pres <- readVECT("aedes_albopictus")
background <- readVECT("background_points")

# 
# Read raster data
#
```

Within R...

```
#  
# Read vector data  
#  
  
# Use sf for vectors  
use_sf()  
  
# Read vector layers  
Aa_pres <- readVECT("aedes_albopictus")  
background <- readVECT("background_points")  
  
#  
# Read raster data  
#  
  
# Use sp for rasters (stars support on the way)
```

Within R...

```
# Read vector layers
Aa_pres <- readVECT("aedes_albopictus")
background <- readVECT("background_points")

#
# Read raster data
#
# 

# Use sp for rasters (stars support on the way)
use_sp()

# List rasters by pattern
worldclim <- execGRASS("g.list",
                         parameters = list(
                           type = "raster",
                           pattern = "worldclim*"))

avσ <- execGRASS("σ.list",
```

Within R...

```
# Use sp for rasters (stars support on the way)
use_sp()

# List rasters by pattern
worldclim <- execGRASS("g.list",
                        parameters = list(
                          type = "raster",
                          pattern = "worldclim*"))

avg <- execGRASS("g.list",
                  parameters = list(
                    type = "raster",
                    pattern = "avg*"))

median <- execGRASS("g.list",
                      parameters = list(
                        type = "raster",
                        pattern = "median*",
                        exclude = "*[1-5]"))
```

Within R...

```
parameters = list(
  type = "raster",
  pattern = "avg*"))

median <- execGRASS("g.list",
  parameters = list(
    type = "raster",
    pattern = "median*",
    exclude = "*[1-5]"))

# Concatenate map lists
to_import <- c(attributes(worldclim)$resOut,
  attributes(avg)$resOut,
  attributes(median)$resOut)

# Read raster layers
predictors <- list()
for (i in to_import){
  predictors[i] <- raster(readRAST(i))
}
```

Within R...

```
parameters = list(  
    type = "raster",  
    pattern = "median*",  
    exclude = "*[1-5]"))  
  
# Concatenate map lists  
to_import <- c(attributes(worldclim)$resOut,  
                  attributes(avg)$resOut,  
                  attributes(median)$resOut)  
  
# Read raster layers  
predictors <- list()  
for (i in to_import){  
    predictors[i] <- raster(readRAST(i))  
}  
  
# Quick visualization in mapview  
mapview(predictors[[3]]) + Aa_pres  
  
#
```

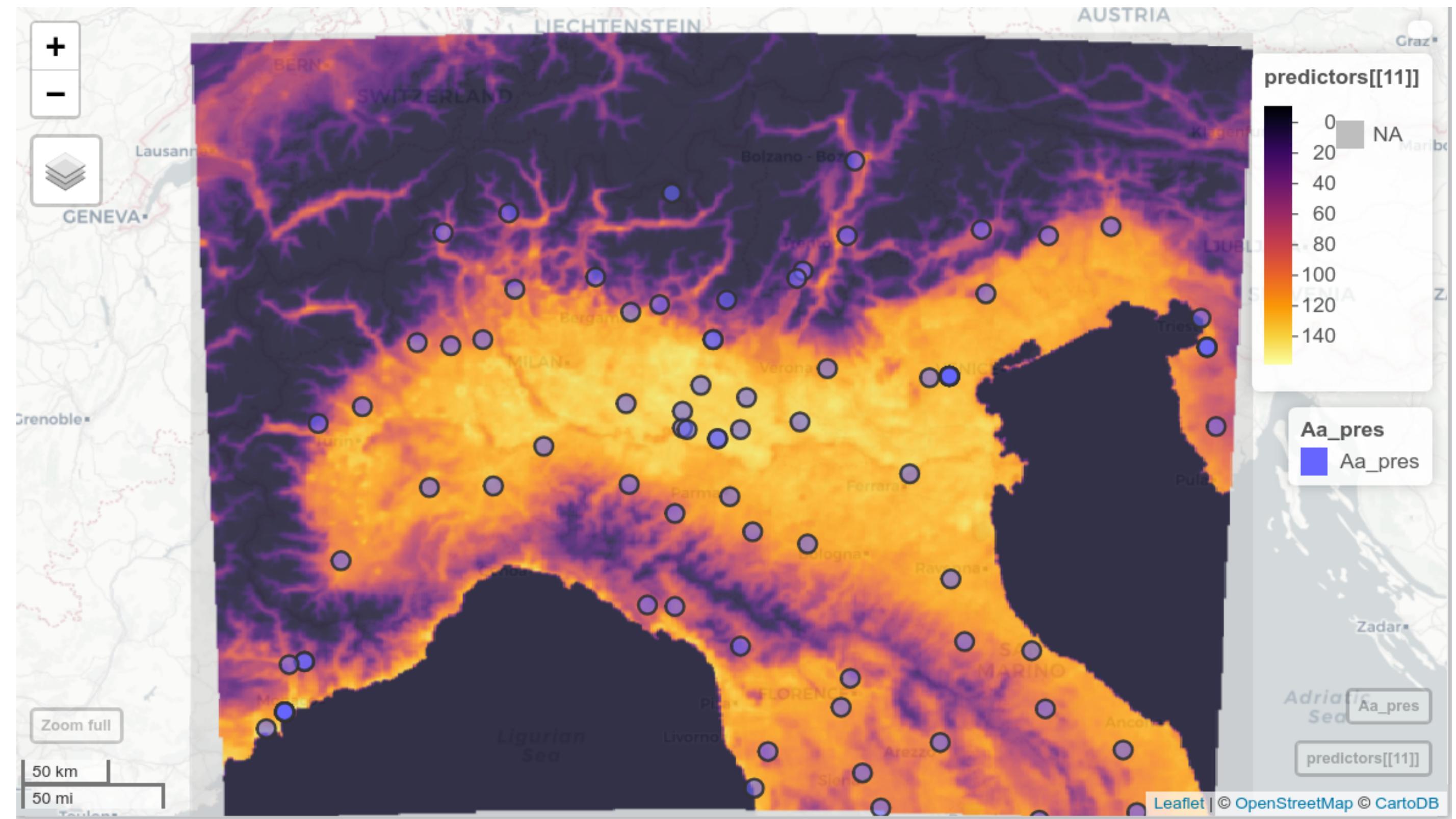
Within R...

```
# Concatenate map lists
to_import <- c(attributes(worldclim)$resOut,
                  attributes(avg)$resOut,
                  attributes(median)$resOut)

# Read raster layers
predictors <- list()
for (i in to_import){
  predictors[i] <- raster(readRAST(i))
}

# Quick visualization in mapview
mapview(predictors[[3]]) + Aa_pres

#
# Data preparation and formatting
#
```



Data formatting

```
#####
# R commands for the session:
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#
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#
# Date: October, 2018
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#####
#
# Install and load required packages
#
```

Data formatting

```
#  
# Data preparation and formatting  
  
# Response variable  
n_pres <- dim(Aa_pres)[1]  
n_backg <- dim(background)[1]  
  
myRespName <- 'Aedes.albopictus'  
  
pres <- rep(1, n_pres)  
backg <- rep(0, n_backg)  
myResp <- c(pres, backg)  
  
myRespXY <- rbind(st_coordinates(Aa_pres),  
                  st_coordinates(background))
```

Data formatting

```
myRespName <- 'Aedes.albopictus'

pres <- rep(1, n_pres)
backg <- rep(0, n_backg)
myResp <- c(pres, backg)

myRespXY <- rbind(st_coordinates(Aa_pres),
                   st_coordinates(background))

# Explanatory variables

# Stack raster layers
myExpl <- raster::stack(predictors)

# All together
myBiomodData <- BIOMOD_FormattingData(resp.var = myResp,
                                         expl.var = myExpl,
                                         resp.xy = myRespXY,
```

Data formatting

```
myRespXY <- rbind(st_coordinates(Aa_pres),  
                   st_coordinates(background))  
  
# Explanatory variables  
  
# Stack raster layers  
myExpl <- raster::stack(predictors)  
  
# All together  
myBiomodData <- BIOMOD_FormattingData(resp.var = myResp,  
                                         expl.var = myExpl,  
                                         resp.xy = myRespXY,  
                                         resp.name = myRespName)  
  
#  
# Run model  
#
```

MaxEnt model

```
#####
# R commands for the session:
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#
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#
# Date: October, 2018
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#####
#
# Install and load required packages
#
```

MaxEnt model

```
exp.var = myExp,
resp.xy = myRespXY,
resp.name = myRespName)

#
# Run model
#



# Set options
myBiomodOption <- BIOMOD_ModelingOptions(
  MAXENT.Phillips =
    list( path_to_maxent.jar = "/home/veroandreo/software/maxent/maxent.jar",
          lq2lqptthreshold = 100,
          l2lqthreshold = 100))

# Run model
myBiomodModelOut <- BIOMOD_Modeling(
  myBiomodData,
  models = c('MAXENT.Phillips'))
```

MaxEnt model

```
# Set options
myBiomodOption <- BIOMOD_ModelingOptions(
  MAXENT.Phillips =
    list( path_to_maxent.jar = "/home/veroandreo/software/maxent/maxent.jar",
          lq2lqptthreshold = 100,
          l2lqthreshold = 100))

# Run model
myBiomodModelOut <- BIOMOD_Modeling(
  myBiomodData,
  models = c('MAXENT.Phillips'),
  models.options = myBiomodOption,
  NbRunEval=5,
  DataSplit=80,
  VarImport=10,
  models.eval.meth = c('ROC','ACCURACY'),
  SaveObj = TRUE,
  rescal.all.models = FALSE,
  do.full.models = FALSE,
  modeling.id = paste(myRespName,"FirstModeling",sep="_"))
```

MaxEnt model

```
myBiomodModelOut <- BIOMOD_Modeling(  
  myBiomodData,  
  models = c('MAXENT.Phillips'),  
  models.options = myBiomodOption,  
  NbRunEval=5,  
  DataSplit=80,  
  VarImport=10,  
  models.eval.meth = c('ROC','ACCURACY'),  
  SaveObj = TRUE,  
  rescal.all.models = FALSE,  
  do.full.models = FALSE,  
  modeling.id = paste(myRespName,"FirstModeling",sep="_"))  
  
# Inspect the model  
myBiomodModelOut  
  
#  
# Model evaluation  
#
```

MaxEnt model

```
models.eval.meth = c('ROC','ACCURACY'),
SaveObj = TRUE,
rescal.all.models = FALSE,
do.full.models = FALSE,
modeling.id = paste(myRespName,"FirstModeling",sep="_"))

# Inspect the model
myBiomodModelOut

#
# Model evaluation
#

# Extract evaluation data
myBiomodModelEval <- get_evaluations(myBiomodModelOut)

# Accuracy
myBiomodModelEval["ACCURACY","Testing.data",,,]
```

MaxEnt model

```
# Inspect the model
myBiomodModelOut

#
# Model evaluation
#
# Extract evaluation data
myBiomodModelEval <- get_evaluations(myBiomodModelOut)

# Accuracy
myBiomodModelEval["ACCURACY","Testing.data",,,]

# ROC: Receiver-operator curve
myBiomodModelEval["ROC","Testing.data",,,]

# Variable importance
get_variables_importance(myBiomodModelOut)
```

MaxEnt model

```
#  
# Model evaluation  
#  
  
# Extract evaluation data  
myBiomodModelEval <- get_evaluations(myBiomodModelOut)  
  
# Accuracy  
myBiomodModelEval["ACCURACY","Testing.data", , , ]  
  
# ROC: Receiver-operator curve  
myBiomodModelEval["ROC","Testing.data", , , ]  
  
# Variable importance  
get_variables_importance(myBiomodModelOut)
```

MaxEnt model

```
# Model evaluation
#
# Extract evaluation data
myBiomodModelEval <- get_evaluations(myBiomodModelOut)

# Accuracy
myBiomodModelEval["ACCURACY","Testing.data",,]

# ROC: Receiver-operator curve
myBiomodModelEval["ROC","Testing.data",,]

# Variable importance
get_variables_importance(myBiomodModelOut)

#
# Model predictions
#
```

Model predictions

```
#####
# R commands for the session:
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#
# Install and load required packages
#
```

Model predictions

```
# Variable importance  
get_variables_importance(myBiomodModelOut)  
  
#  
# Model predictions  
#  
  
# Set parameters for model projection  
myBiomodProj <- BIOMOD_Projection(  
    modeling.output = myBiomodModelOut,  
    new.env = myExpl,  
    proj.name = "current",  
    selected.models = "all",  
    compress = FALSE,  
    build.clamping.mask = FALSE)  
  
# Obtain predictions  
mod_proj <- get_predictions(myBiomodProj)
```

Model predictions

```
# Model predictions
#
#
# Set parameters for model projection
myBiomodProj <- BIOMOD_Projection(
  modeling.output = myBiomodModelOut,
  new.env = myExpl,
  proj.name = "current",
  selected.models = "all",
  compress = FALSE,
  build.clamping.mask = FALSE)

# Obtain predictions
mod_proj <- get_predictions(myBiomodProj)
mod_proj

# Plot predicted model
plot(mod_proj)
plot(mod_proj[[2]], main = "Predicted potential distribution")
```

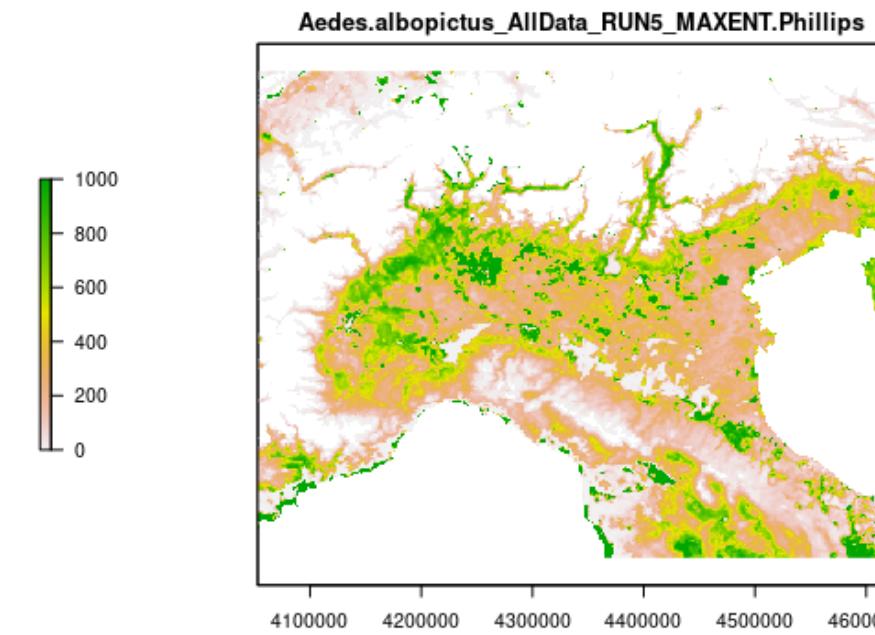
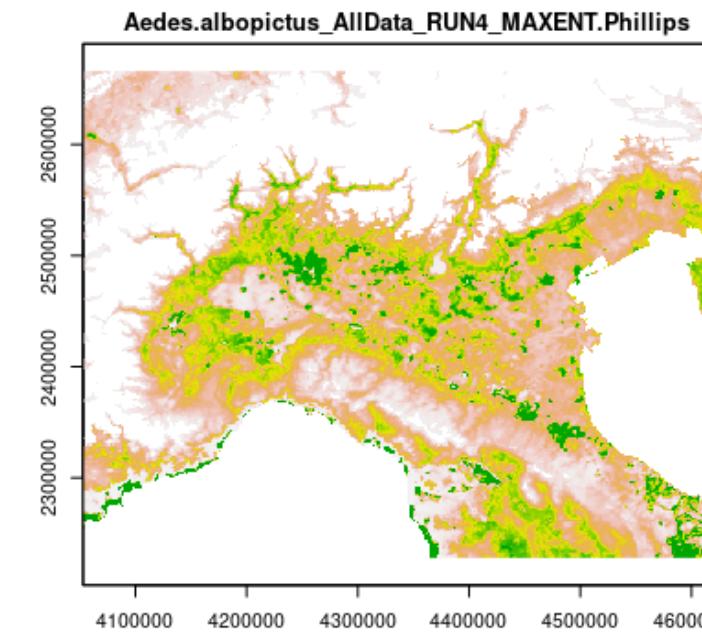
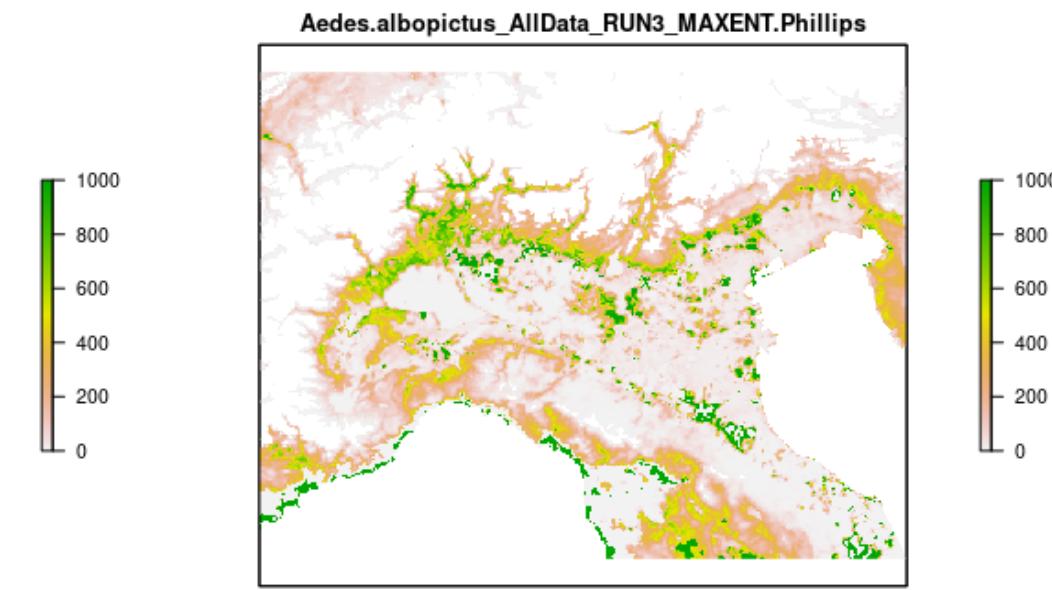
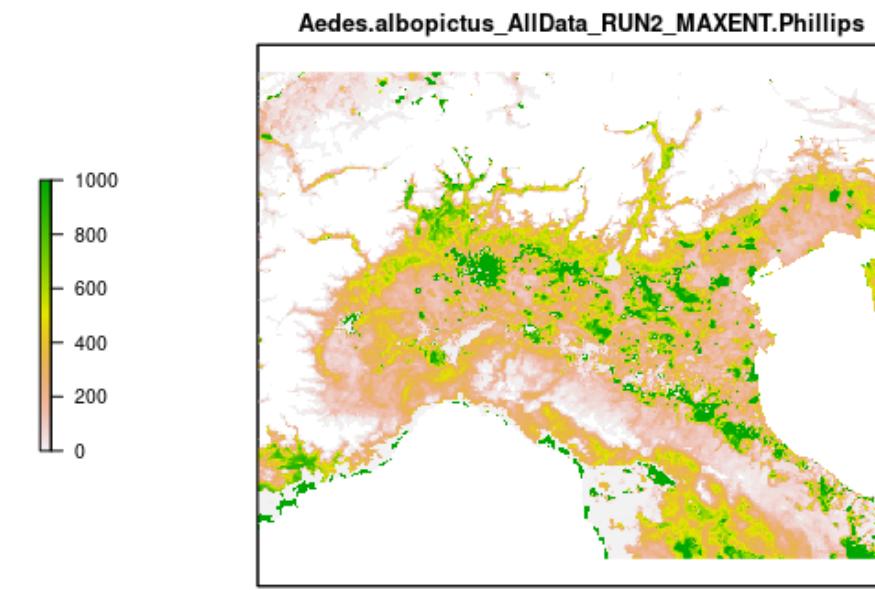
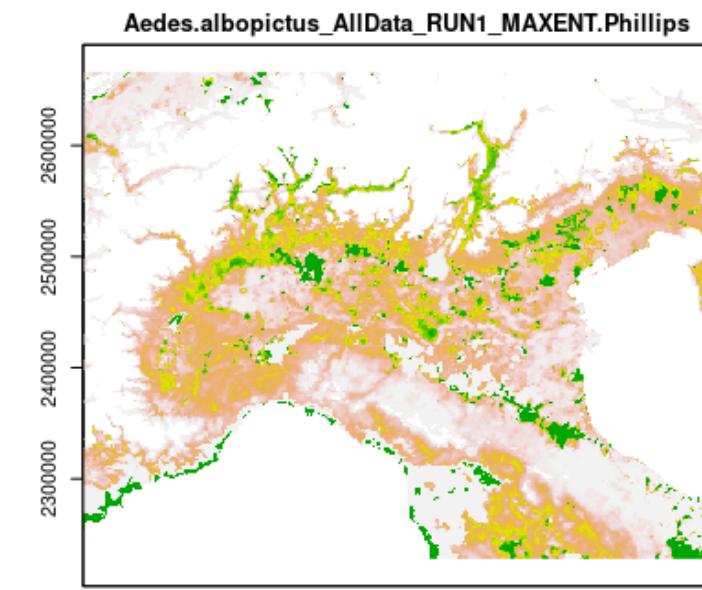
Model predictions

```
# Set parameters for model projection
myBiomodProj <- BIOMOD_Projection(
  modeling.output = myBiomodModelOut,
  new.env = myExpl,
  proj.name = "current",
  selected.models = "all",
  compress = FALSE,
  build.clamping.mask = FALSE)

# Obtain predictions
mod_proj <- get_predictions(myBiomodProj)
mod_proj

# Plot predicted model
plot(mod_proj)
plot(mod_proj[[2]], main = "Predicted potential distribution")

#
# Optionally, write data back to GRASS GIS
```





Task: Explore the algorithms available in `BIOMOD_Modeling()` and try with a different one. Compare assumptions and results.

Write model predictions into GRASS GIS DB

```
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#
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#
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# Updated: August, 2019
#####
#
# Install and load required packages
#
```

Write model predictions into GRASS GIS DB

```
# Obtain predictions
mod_proj <- get_predictions(myBiomodProj)
mod_proj

# Plot predicted model
plot(mod_proj)
plot(mod_proj[[2]], main = "Predicted potential distribution")

#
# Optionally, write data back to GRASS GIS
# 

# Only one layer
g <- as(mod_proj[[4]], 'SpatialGridDataFrame')
writeRAST(g, "maxent_albopictus", flags = "overwrite")

# Export all MaxEnt runs
# ...
```

Write model predictions into GRASS GIS DB

```
# Obtain predictions
mod_proj <- get_predictions(myBiomodProj)
mod_proj

# Plot predicted model
plot(mod_proj)
plot(mod_proj[[2]], main = "Predicted potential distribution")

#
# Optionally, write data back to GRASS GIS
#
# Only one layer
g <- as(mod_proj[[4]], 'SpatialGridDataFrame')
writeRAST(g, "maxent_albopictus", flags = "overwrite")

# Export all MaxEnt runs
# writeRaster(mod_proj, "maxent_all", flags = "overwrite")
```

QUESTIONS?



Other (very) useful resources

- Temporal data processing wiki
- GRASS GIS and R for time series processing wiki
- GRASS GIS temporal workshop at NCSU
- GRASS GIS workshop held in Jena 2018
- GRASS GIS course IRSAE 2018
- GRASS GIS course in Argentina 2018

References

- Gebbert, S., Pebesma, E. (2014). *A temporal GIS for field based environmental modeling*. Environmental Modelling & Software, 53, 1-12. [DOI](#)
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- Gebbert, S., Leppelt, T. and Pebesma, E. (2019). *A Topology Based Spatio-Temporal Map Algebra for Big Data Analysis*. Data, 4, 86. [DOI](#)



Thanks for your attention!!



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