

**NERRS / SWMP**

**Training Workshop: *R* Intro & SWMP*r***

**October 25, 2015**

## SWMP*r* analyze

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# Objectives for the session

- What are some basic analyses that can be accomplished with SWMPr?
- What are some plotting functions provided by SWMPr?
- What are some resources for additional learning?

# Interactive portion

We will use the `swmpr2.Rproj` project for this session, double-click to open in RStudio

- location on flash drive
- location online

## Interactive portion

We will use the `swmpr2.Rproj` project for this session, double-click to open in RStudio

- location on flash drive
- location online

You will run examples whenever you see this guy:



Don't forget to use your stickies: **green** for done/ok, **red** for problem

# Basic analyses with SWMP<sub>r</sub>

How do we want to use the data?

# Basic analyses with SWMP<sub>r</sub>

How do we want to use the data?

- What has happened at my site over time?
- Are there differences between sites?
- Can we remove seasonal trends?
- Are there differences between parameters?
- Others?

# Basic analyses with SWMP<sub>r</sub>

Take a few minutes to acquaint yourself with the *analyze* functions:

```
help.search('analyze', package = 'SWMPr')
```

# Basic analyses with SWMP<sub>r</sub>



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```
help.search('analyze', package = 'SWMPr')
```

Which functions simplify the data?

Which functions could you use to explore or visualize the data?

Which functions are related to metabolism?



# Basic analyses with SWMPr - missing data

Most datasets will have missing values - how do you deal with those?

Remove? Set as mean? Replace with similar?

SWMPr provides a function to interpolate missing data: `na.approx`

To start, let's import and plot some data...

# Basic analyses with SWMP<sub>r</sub> - missing data



- Import the 2012 water quality data for cbmmc from the 'zip\_ex' folder
- Deal with QAQC columns
- Select two columns of interest
- Plot the data subset (`?plot.swmpr`)

# Basic analyses with SWMP<sub>r</sub> - missing data



- Import the 2012 water quality data for cbmmc from the 'zip\_ex' folder

```
mypath <- 'zip_ex'  
dat <- import_local(mypath, 'cbmmcwq2012')
```

- Deal with QAQC columns
- Select two columns of interest
- Plot the data subset (?plot.swmpr)

# Basic analyses with SWMP<sub>r</sub> - missing data



- Import the 2012 water quality data for cbmmc from the 'zip\_ex' folder

```
mypath <- 'zip_ex'  
dat <- import_local(mypath, 'cbmmcwq2012')
```

- Deal with QAQC columns

```
tmp <- qaqc(dat)
```

- Select two columns of interest
- Plot the data subset (?plot.swmpr)

# Basic analyses with SWMP<sub>r</sub> - missing data



- Import the 2012 water quality data for cbmmc from the 'zip\_ex' folder

```
mypath <- 'zip_ex'  
dat <- import_local(mypath, 'cbmmcwq2012')
```

- Deal with QAQC columns

```
tmp <- qaqc(dat)
```

- Select two columns of interest

```
tmp <- subset(tmp, select = 'do_mgl', subset = c('2012-10-01 0:0',  
          '2012-10-31 0:0'))
```

- Plot the data subset (?plot.swmpr)

# Basic analyses with SWMP<sub>r</sub> - missing data



- Import the 2012 water quality data for cbmmc from the 'zip\_ex' folder

```
mypath <- 'zip_ex'  
dat <- import_local(mypath, 'cbmmcwq2012')
```

- Deal with QAQC columns

```
tmp <- qaqc(dat)
```

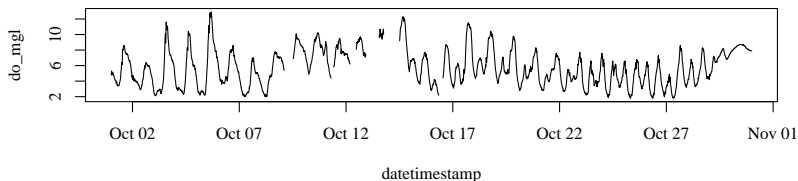
- Select two columns of interest

```
tmp <- subset(tmp, select = 'do_mgl', subset = c('2012-10-01 0:0',  
          '2012-10-31 0:0'))
```

- Plot the data subset (?plot.swmpr)

```
plot(tmp)
```

# Basic analyses with SWMP<sub>r</sub> - missing data



Notice the missing data around October 12<sup>th</sup>

The `na.approx` function (`?na.approx.swmpr`) has three arguments:

- **object**: swmpr data object to fill
- **params**: name(s) of parameter to fill
- **maxgap**: maximum gap size to interpolate

# Basic analyses with SWMP<sub>r</sub> - missing data



- Use `na.approx` to interpolate the missing data (`?na.approx.swmpr`)
- Plot the two to see the differences (`?plot.swmpr`)



# Basic analyses with SWMP<sub>r</sub> - missing data



- Use `na.approx` to interpolate the missing data (`?na.approx.swmpr`)

```
tmp2 <- na.approx(tmp, params = 'do_mgl', maxgap = 100)
```

- Plot the two to see the differences (`?plot.swmpr`)

# Basic analyses with SWMPr - missing data

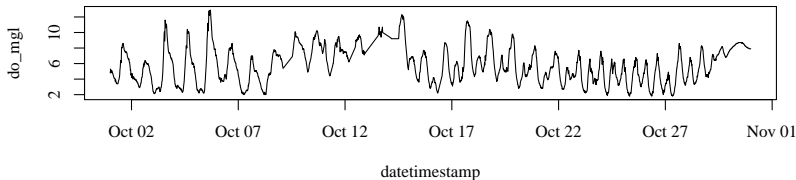
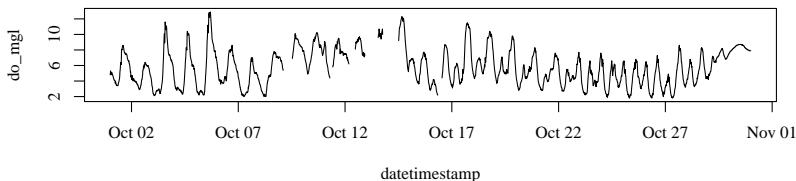


- Use `na.approx` to interpolate the missing data (`?na.approx.swmpr`)

```
tmp2 <- na.approx(tmp, params = 'do_mgl', maxgap = 100)
```

- Plot the two to see the differences (`?plot.swmpr`)

```
plot(tmp)  
plot(tmp2)
```

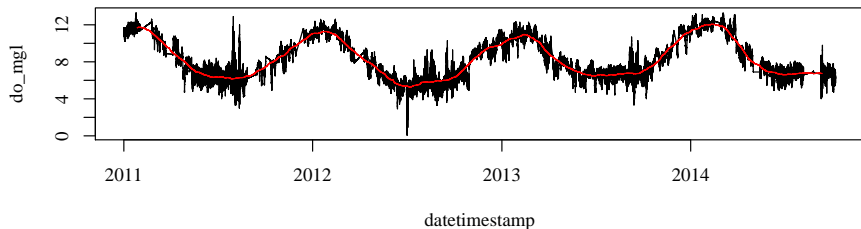


# Basic analyses with SWMP<sub>r</sub> - smoothing data

Now we know how to fill missing data, let's see how it can help...

Trend evaluation often considers variation at different spatial scales

As an example, we want to evaluate variation in dissolved oxygen at seasonal or annual scales



## Analysis 2 - Smoothing and aggregation

The **smoother** function (`?smoother`) calculates a moving window average of a time series

- **x**: Input data object
- **window**: the size of the smoothing window, defaults to five observations at the current time step
- **sides**: what defines the window, centered on an observation (2, default) or use only the preceding observations (1)
- **params**: which parameters to smooth, default all

What would be a good window to look at seasonal or annual variation?

# Basic analyses with SWMP<sub>r</sub> - smoothing data



- Import all years of water quality data for cbmip from the ‘zip\_ex’ folder
- Deal with QAQC columns and subset DO
- Use **smoother** to remove daily and short-term variation, which window to use?
- Plot results (see `?plot.swmpr` and `?lines.swmpr`)

# Basic analyses with SWMP<sub>r</sub> - smoothing data



- Import all years of water quality data for cbmip from the 'zip\_ex' folder

```
mypath <- 'zip_ex'  
dat <- import_local(mypath, 'cbmipwq')
```

- Deal with QAQC columns and subset DO
- Use **smoother** to remove daily and short-term variation, which window to use?
- Plot results (see ?plot.swmpr and ?lines.swmpr)

# Basic analyses with SWMP<sub>r</sub> - smoothing data



- Import all years of water quality data for cbmip from the 'zip\_ex' folder

```
mypath <- 'zip_ex'  
dat <- import_local(mypath, 'cbmipwq')
```

- Deal with QAQC columns and subset DO

```
tmp <- qaqc(dat)  
tmp <- subset(dat, select = 'do_mgl')
```

- Use **smoother** to remove daily and short-term variation, which window to use?
- Plot results (see ?plot.swmpr and ?lines.swmpr)

# Basic analyses with SWMP<sub>r</sub> - smoothing data



- Import all years of water quality data for cbmip from the 'zip\_ex' folder

```
mypath <- 'zip_ex'  
dat <- import_local(mypath, 'cbmipwq')
```

- Deal with QAQC columns and subset DO

```
tmp <- qaqc(dat)  
tmp <- subset(dat, select = 'do_mgl')
```

- Use **smoother** to remove daily and short-term variation, which window to use?

```
do_smooth <- smoother(tmp2, window = 5000)
```

- Plot results (see ?plot.swmpr and ?lines.swmpr)



# Basic analyses with SWMP<sub>r</sub> - smoothing data



- Import all years of water quality data for cbmip from the 'zip\_ex' folder

```
mypath <- 'zip_ex'  
dat <- import_local(mypath, 'cbmipwq')
```

- Deal with QAQC columns and subset DO

```
tmp <- qaqc(dat)  
tmp <- subset(dat, select = 'do_mgl')
```

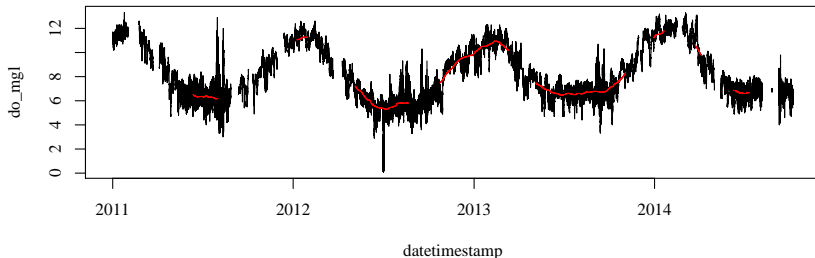
- Use **smoother** to remove daily and short-term variation, which window to use?

```
do_smooth <- smoother(tmp2, window = 5000)
```

- Plot results (see ?plot.swmpr and ?lines.swmpr)

```
plot(tmp)  
lines(do_smooth)
```

# Basic analyses with SWMPr - smoothing data



What happened?

How can we fix the problem?

# Basic analyses with SWMPr - smoothing data



Repeat the analysis but use `na.approx` to fill missing data

```
mypath <- 'zip_ex'
dat <- import_local(mypath, 'cbmipwq')
tmp <- qaqc(dat)
tmp <- subset(tmp, select = 'do_mgl')
do_smooth <- smoother(tmp, window = 5000)
plot(tmp)
lines(do_smooth, col = 'red')
```

Where would you use `na.approx`?

# Basic analyses with SWMP<sub>r</sub> - smoothing data



Repeat the analysis but use `na.approx` to fill missing data

```
mypath <- 'zip_ex'
dat <- import_local(mypath, 'cbmipwq')
tmp <- qaqc(dat)
tmp <- subset(tmp, select = 'do_mgl')
do_smooth <- smoother(tmp, window = 5000)
plot(tmp)
lines(do_smooth, col = 'red')
```

Where would you use `na.approx`?

```
mypath <- 'zip_ex'
dat <- import_local(mypath, 'cbmipwq')
tmp <- qaqc(dat)
tmp <- subset(tmp, select = 'do_mgl')
tmp <- na.approx(tmp, maxgap = 5000)
do_smooth <- smoother(tmp, window = 5000)
plot(tmp)
lines(do_smooth, col = 'red')
```

# Basic analyses with SWMPr - smoothing data



Repeat the analysis but use `na.approx` to fill missing data

```
mypath <- 'zip_ex'
dat <- import_local(mypath, 'cbmipwq')
tmp <- qaqc(dat)
tmp <- subset(tmp, select = 'do_mgl')
do_smooth <- smoother(tmp, window = 5000)
plot(tmp)
lines(do_smooth, col = 'red')
```

Where would you use `na.approx`?

```
mypath <- 'zip_ex'
dat <- import_local(mypath, 'cbmipwq')
tmp <- qaqc(dat)
tmp <- subset(tmp, select = 'do_mgl')
tmp <- na.approx(tmp, maxgap = 5000)
do_smooth <- smoother(tmp, window = 5000)
plot(tmp)
lines(do_smooth, col = 'red')
```

Bonus: Try changing `maxgap` or `window`

# Basic analyses with SWMPr - aggregating data

Finally, we can use `aggreswmp` to summarize and plot for an alternative interpretation

`aggreswmp` has five main arguments:

- `swmpr_in`: input data object
- `by`: aggregation period ('years', 'quarters', etc.)
- `FUN`: aggregation function, defaults to mean
- `params`: which parameters to aggregate, defaults all
- `aggs_out`: get the raw data, use this to make plots

# Basic analyses with SWMP<sub>r</sub> - aggregating data



- Import all years of water quality data for cbmip from the 'zip\_ex' folder, QAQC cleanup, and subset DO
- Use `aggreswmp` (`?aggreswmp`) to get quarterly summaries of the data

# Basic analyses with SWMPr - aggregating data



- Import all years of water quality data for cbmip from the 'zip\_ex' folder, QAQC cleanup, and subset DO

```
mypath <- 'zip_ex'
dat <- import_local(mypath, 'cbmipwq')
tmp <- qaqc(dat)
tmp <- subset(dat, select = 'do_mgl')
```

- Use `aggreswmp` (?aggreswmp) to get quarterly summaries of the data



# Basic analyses with SWMP<sup>r</sup> - aggregating data



- Import all years of water quality data for cbmip from the 'zip\_ex' folder, QAQC cleanup, and subset DO

```
mypath <- 'zip_ex'
dat <- import_local(mypath, 'cbmipwq')
tmp <- qaqc(dat)
tmp <- subset(dat, select = 'do_mgl')
```

- Use `aggreswmp` (?aggreswmp) to get quarterly summaries of the data

```
aggtmp <- aggreswmp(tmp, by = 'quarters')
```

# Basic analyses with SWMP<sub>r</sub> - aggregating data



- Import all years of water quality data for cbmip from the 'zip\_ex' folder, QAQC cleanup, and subset DO

```
mypath <- 'zip_ex'
dat <- import_local(mypath, 'cbmipwq')
tmp <- qaqc(dat)
tmp <- subset(dat, select = 'do_mgl')
```

- Use `aggreswmp` (`?aggreswmp`) to get quarterly summaries of the data

```
aggtmp <- aggreswmp(tmp, by = 'quarters')
```

- Bonus: Try different aggregation periods
- Bonus: Try different aggregation functions

# Basic analyses with SWMPr - aggregating data



- Import all years of water quality data for cbmip from the 'zip\_ex' folder, QAQC cleanup, and subset DO

```
mypath <- 'zip_ex'
dat <- import_local(mypath, 'cbmipwq')
tmp <- qaqc(dat)
tmp <- subset(dat, select = 'do_mgl')
```

- Use `aggreswmp` (`?aggreswmp`) to get quarterly summaries of the data

```
aggtmp <- aggreswmp(tmp, by = 'quarters')
```

- Bonus: Try different aggregation periods

```
aggtmp <- aggreswmp(tmp, by = 'years')
aggtmp <- aggreswmp(tmp, by = 'weeks')
```

- Bonus: Try different aggregation functions

# Basic analyses with SWMPr - aggregating data



- Import all years of water quality data for cbmip from the 'zip\_ex' folder, QAQC cleanup, and subset DO

```
mypath <- 'zip_ex'  
dat <- import_local(mypath, 'cbmipwq')  
tmp <- qaqc(dat)  
tmp <- subset(dat, select = 'do_mgl')
```

- Use `aggreswmp` (?`aggreswmp`) to get quarterly summaries of the data

```
aggtmp <- aggreswmp(tmp, by = 'quarters')
```

- Bonus: Try different aggregation periods

```
aggtmp <- aggreswmp(tmp, by = 'years')  
aggtmp <- aggreswmp(tmp, by = 'weeks')
```

- Bonus: Try different aggregation functions

```
fun_in <- function(x) var(x, na.rm = TRUE)  
aggtmp <- aggreswmp(swmpr_in, FUN = fun_in, 'years')
```

# Basic analyses with SWMP<sub>r</sub> - aggregating data



Plot the aggregated data by quarters - use `aggs_out = TRUE`

# Basic analyses with SWMP<sub>r</sub> - aggregating data



Plot the aggregated data by quarters - use `aggs_out = TRUE`

```
# use aggs_out to get all  
aggtmp <- aggreswmp(tmp, by = 'quarters', aggs_out = TRUE)
```

# Basic analyses with SWMPr - aggregating data



Plot the aggregated data by quarters - use `aggs_out = TRUE`

```
# use aggs_out to get all  
aggtmp <- aggreswmp(tmp, by = 'quarters', aggs_out = TRUE)
```

Then use `boxplot` (`?boxplot`) from the R stats package

# Basic analyses with SWMPPr - aggregating data

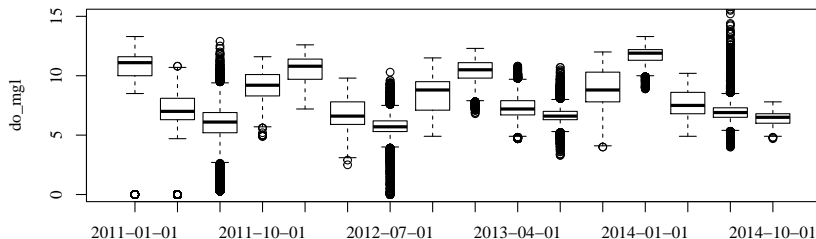


Plot the aggregated data by quarters - use `aggs_out = TRUE`

```
# use aggs_out to get all  
aggtmp <- aggreswmp(tmp, by = 'quarters', aggs_out = TRUE)
```

Then use `boxplot` (`?boxplot`) from the R stats package

```
# use boxplot  
boxplot(do_mgl ~ datetimestamp, data = aggtmp, ylab = 'do_mgl', ylim = c(0, 15))
```





# Basic analyses with SWMPPr - aggregating data

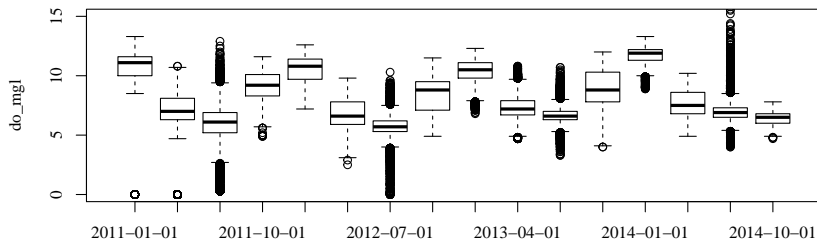


Plot the aggregated data by quarters - use `aggs_out = TRUE`

```
# use aggs_out to get all  
aggtmp <- aggreswmp(tmp, by = 'quarters', aggs_out = TRUE)
```

Then use `boxplot` (`?boxplot`) from the R stats package

```
# use boxplot  
boxplot(do_mgl ~ datetimestamp, data = aggtmp, ylab = 'do_mgl', ylim = c(0, 15))
```



Have a look at the data from `aggs_out = TRUE` and `aggs_out = FALSE`, how do they differ and why?

# Plotting functions in SWMP<sub>r</sub>

R provides near limitless options to visualize data - a full coverage of these tools would take days

We will briefly go over some key plotting functions in SWMP<sub>r</sub>, each is designed for simplicity and efficiency to summarize lots of data

Plotting functions in SWMP<sub>r</sub>:

- `decomp`: time series decomposition
- `decomp_cj`: time series decomposition, alternative
- `map_reserve`: plot a basic map of a reserve
- `overplot`: plot multiple parameters on the same plot
- `plot_metab`: plot metabolism estimates
- `plot_summary`: plot multiple summaries for a parameter

# Plotting functions in SWMP<sub>r</sub>

The `map_reserve` function can be used to map sites:

- `nerr_site_id`: site(s) to map, usually first three letters
- `zoom`: zoom factor for the map (usually between 5–15)
- `map_type`: 'terrain', 'satellite', 'roadmap', or 'hybrid'

```
# try any of these examples  
map_reserve('jac')  
  
map_reserve('elk', zoom = 13,  
            map_type = 'hybrid')  
  
map_reserve('gtmss', zoom = 15,  
            map_type = 'satellite',  
            text_col = 'lightblue')
```

# Plotting functions in SWMP<sup>r</sup>

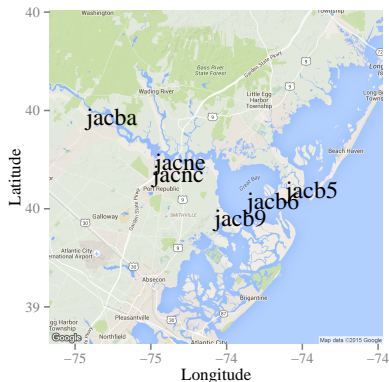
The `map_reserve` function can be used to map sites:

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# try any of these examples
map_reserve('jac')

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  map_type = 'hybrid')

map_reserve('gtmss', zoom = 15,
  map_type = 'satellite',
  text_col = 'lightblue')
```



# Plotting functions in SWMP<sub>r</sub>

The `overplot` function can be used to plot one to many time series:

- `dat_in`: input `swmpr` object
- `select`: parameter(s) to plot (passed to `subset`)
- `subset`: date ranges to plot (passed to `subset`)

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Import the 2011 data for `cbmip`

Plot DO, temperature, and salinity for August (`?overplot`)

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- `dat_in`: input `swmpr` object
- `select`: parameter(s) to plot (passed to `subset`)
- `subset`: date ranges to plot (passed to `subset`)

Import the 2011 data for cbmip

```
# import  
mypath <- 'zip_ex'  
dat <- import_local(mypath, 'cbmipwq2011')
```

Plot DO, temperature, and salinity for August (?`overplot`)

# Plotting functions in SWMP<sup>r</sup>

The `overplot` function can be used to plot one to many time series:

- `dat_in`: input `swmpr` object
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Import the 2011 data for `cbmip`

```
# import  
mypath <- 'zip_ex'  
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```

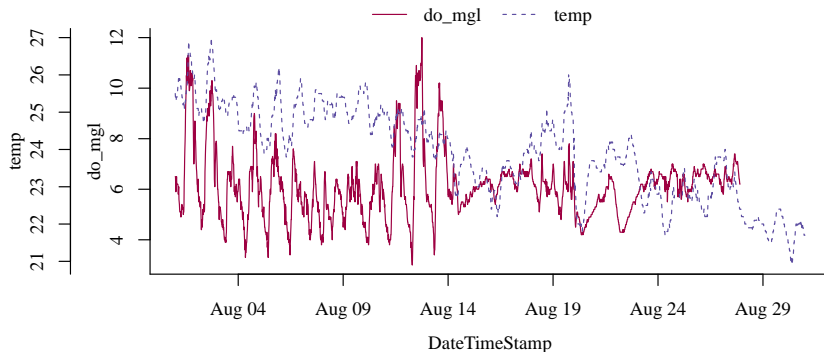
Plot DO, temperature, and salinity for August (?`overplot`)

```
# plot  
overplot(dat, select = c('do_mgl', 'temp'),  
  subset = c('2011-08-01 0:0', '2011-08-31 0:0'))
```



# Plotting functions in SWMP<sub>r</sub>

```
# plot  
overplot(dat, select = c('do_mgl', 'temp'),  
  subset = c('2011-08-01 0:0', '2011-08-31 0:0'))
```



# Plotting functions in SWMP<sup>r</sup>

The `plot_summary` function can be used to summarize a parameter across the time series

- `swmpr_in`: input `swmpr` object
- `param`: parameter to summarize
- `years`: years to plot

Import all years of data for `cbmip`, plot a summary of temperature (`?plot_summary`)

# Plotting functions in SWMP<sup>r</sup>

The `plot_summary` function can be used to summarize a parameter across the time series

- `swmpr_in`: input `swmpr` object
- `param`: parameter to summarize
- `years`: years to plot

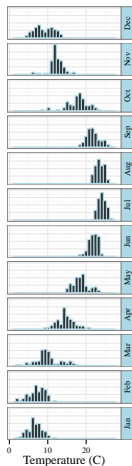
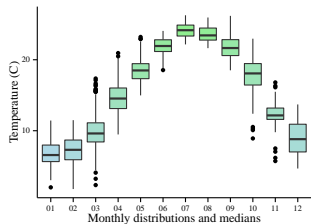
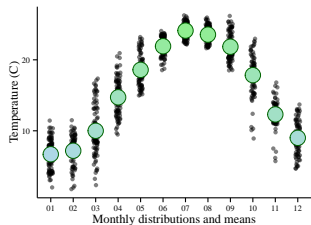
Import all years of data for `cbmip`, plot a summary of temperature (`?plot_summary`)

```
# import
mypath <- 'zip_ex'
dat <- import_local(mypath, 'cbmipwq')

# plot
plot_summary(dat, 'temp')
```

# Plotting functions in SWMP<sub>r</sub>

```
# plot  
plot_summary(dat, 'temp')
```



## Additional resources

You've been exposed to some basic tools to *retrieve*, *organize*, and *analyze* SWMP data with SWMP<sub>r</sub>

There are multiple resources available for continued learning:

- [swmprats.net](http://swmprats.net)
  - ▶ User forum to post questions, plot of the month
  - ▶ Widgets for data viz
  - ▶ Access to workshop materials
  - ▶ Access to 2014 workshop materials
- SWMP<sub>r</sub> online reference [manual](#): list of all functions
- SWMP<sub>r</sub> cookbook: step-by-step scripts as stand-alone analyses
- Email instructors: [beck.marcus@epa.gov](mailto:beck.marcus@epa.gov), [todd.obrien@noaa.gov](mailto:todd.obrien@noaa.gov)

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*Questions??*