# R Scripts for Eddy Covariance and Storage Flux Systems

Date: 2016-01-19 (Tuesday)

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### Preface

- Workshop requirements
  - Attended the Statistical Analyses using R course.
  - PC or Mac installed with R (and RStudio).
  - Internet
- Workshop schedule (tentative)
  - 9 10 AM: PC/Mac basic setup (*Version control* GitHub; R and RStudio)
  - 10 1 PM: Data analysis of Eddy Covariance and Biomet Systems (including tea break if any)
  - 1 2 PM: Lunch
  - 2 3 PM: Data analysis of Storage Flux systems
  - 3 4 PM: Data analysis of Soil Chamber systems (including tea break if any)

### Workshop Outcomes

- By the end of this workshop, you will be able to,
  - Understand the developed R script.
  - Use the R scripts.
  - Modify the R scripts.

## PC/Mac Basic Setup



- Make sure you have R and RStudio installed.
- Now, we are going to download the scripts stored in an online repository called GitHub (a version control software system).
  - Go to the website: <a href="http://github.com/yusriy/fluxMPOB">http://github.com/yusriy/fluxMPOB</a>
  - Download the ZIP file by clicking the *Download ZIP* button.
  - Unzip and place the downloaded folder *fluxMPOB-master* into your *Documents* folder.
  - Double-click the *fluxMPOB.Rproj* file, RStudio should run.

#### R Folder Structure

- Within the R Project folder there are sub-folders:
  - Data
    - Where you should store your data.
  - R
    - Where you should keep your R scripts.
  - Extra: Figs
    - Where the output in the form of figures should be written to.
  - Docs
    - Where you should keep the relevant documents or manuscripts.
  - README.md
    - A *Markdown readme* file to explain about the Project to users. Recommended to have for every project, especially in GitHub.

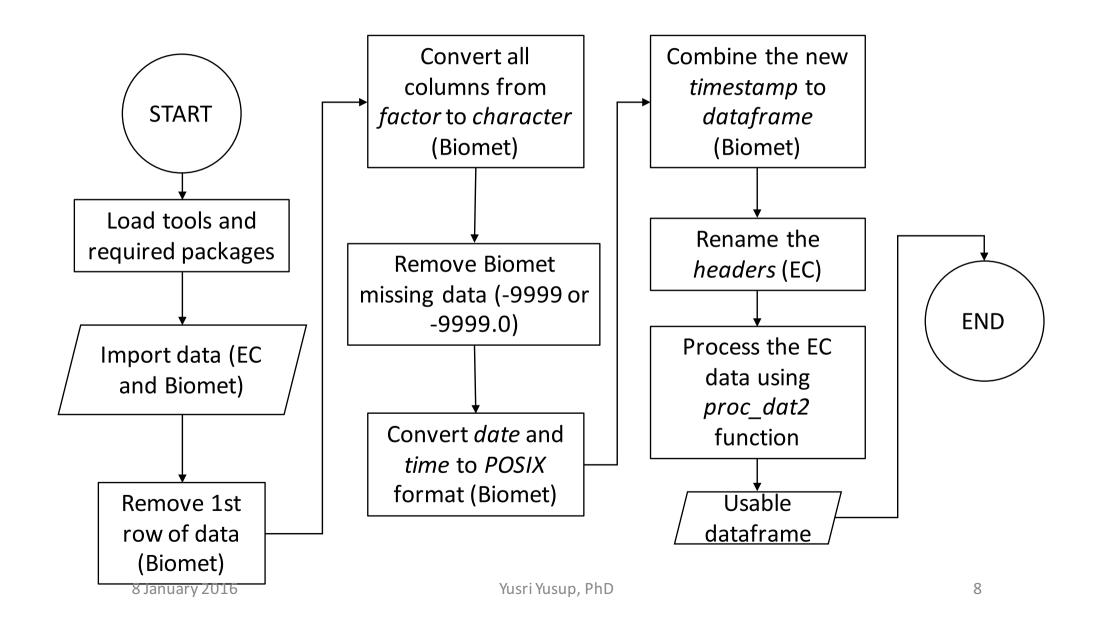
# Eddy Covariance and Biomet Systems

- Output from EddyPro
  - Within (usually) RESULTS folder of the Result\_EDDYPRO folder.
  - EC Full Output contains all measured and calculated parameters of EC (many of them you will not use)
  - Meteorological data Biomet contains only meteorological data.
  - The timestamp of both EC and Biomet should be the same. – the function of the Sutron datalogger.
    - Very important because this is how we can merge both datasets together.

### EC and Biomet Parameters

- Open the Full Output data file in the data folder.
- A <u>LOT</u> of parameters/variables with many of them unimportant but I recommend using this dataset instead of the *Essential* dataset because you never know when you might want to refer to a any of the variables for troubleshooting, e.g., qc (quality control), etc.
- Now, we discuss on some of the variables in detail.
- ... and **study** the EC data import and processing R script.

### EC and Biomet R Script Flowchart



# Understanding the EC and Biomet data

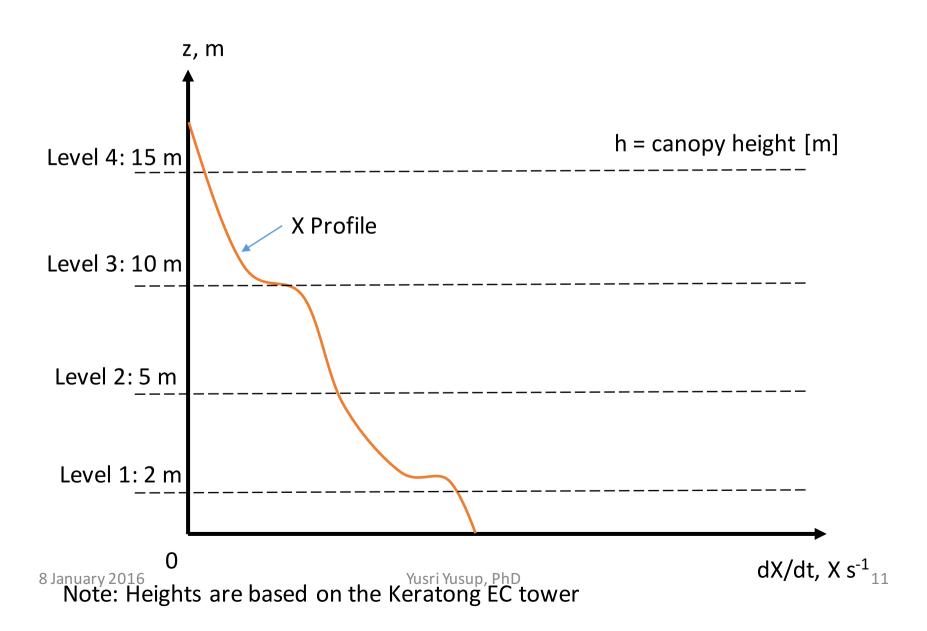
• Start with summary(), plot(), hist(), etc.

# Storage Fluxes (LE, H, CO2) Parameters (Profiler)

- Storage flux equations for LE, H, and CO<sub>2</sub> \*(Finnigan, 2006) used are the same as the ones employed by LI-COR.
- The units of LE, H, and CO<sub>2</sub> are the same as given by EC [W m<sup>-2</sup>].
- Multi-level measurements of H<sub>2</sub>O, T, and CO<sub>2</sub> are needed to calculated amount of energy/concentration flux stored in the canopy.
- H and LE storage fluxes can also be calculated from the EC dataset since multi-level T and RH were also collected.

<sup>\*</sup>Finnigan, J. (2006). The storage term in eddy flux calculations. *Agricultural and Forest Meteorology*. 136, 108-113. Yusri Yusup, PhD

### Multi-level Measurements



### Storage Flux Equations

LE storage flux = 
$$\frac{\lambda}{MW} \int_0^h \frac{dH_2O}{dt} dz$$
 [W m<sup>-2</sup>]

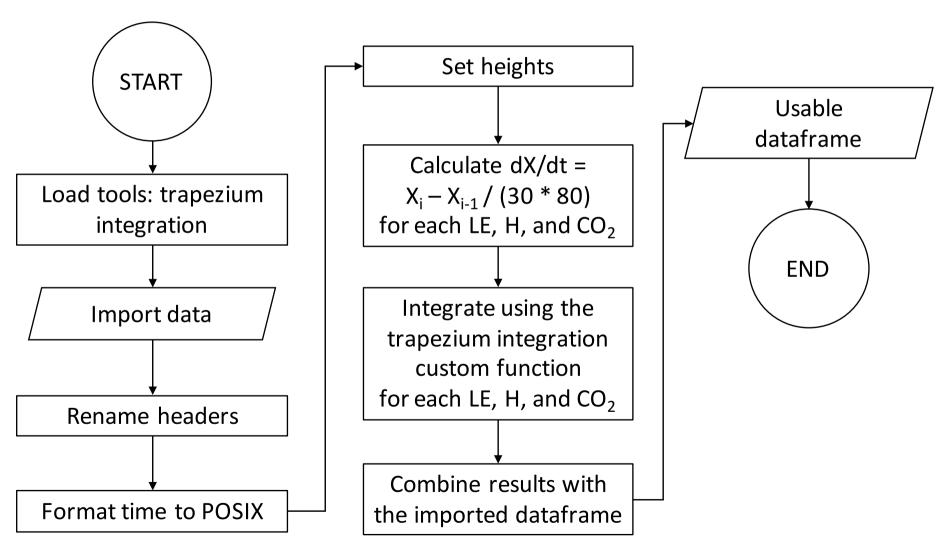
 $\lambda$  = latent heat of vaporization = 2540000 J kg<sup>-1</sup> MW = 18 kg mol H<sub>2</sub>O<sup>-1</sup>

$$H storage flux = \rho_{dry air} c_{p,dry air} \int_{0}^{h} \frac{dT}{dt} dz \qquad [W m^{-2}]$$

hodry air = density of dry air = 1.2754 kg m-3 c<sub>p, dry air</sub> = heat capacity of dry air = 1005 J kg<sup>-1</sup> K<sup>-1</sup>

$$CO_2 storage flux = \int_0^h \frac{dCO_2}{dt} dz$$
 [µmol m<sup>-2</sup> s<sup>-1</sup>]

### Storage Flux R Script Flowchart

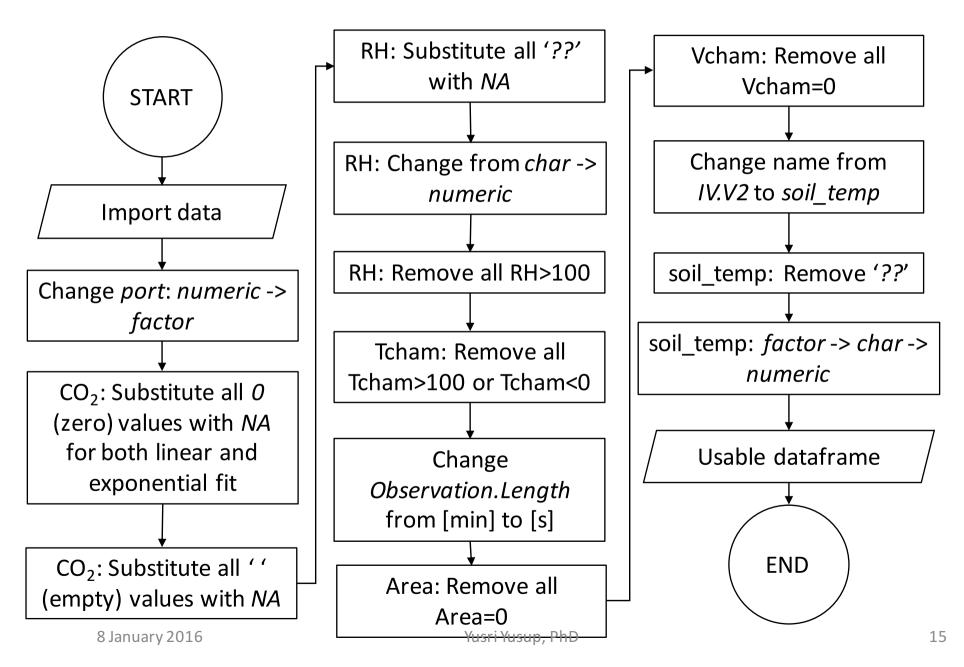


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#### Soil Chamber Flux Parameters

- Different sampling points are connected to different ports of the analyzer (Multiplexer).
- Four soil flux chambers:
  - Port 1: Harvest path under palm
  - Port 2: Harvest path open
  - Port 3: Frond pile north
  - Port 4: Frond pile south
  - Port 5 8: Profiler (not used here since the data is added to the storage flux dataset)
- Again, not all parameters/columns will be used.
- Data that we import into R is assumed to be already processed by LI-COR's SoilFluxPro.

### Soil Chamber Flux R Script Flowchart



### END