```
title: "FLUXNET-CH4 Upscaling"
author: "Gavin McNicol"
date: "2/22/2021"
output: html_document
```{r setup, include=FALSE}
knitr::opts_chunk$set(echo = TRUE)
Load packages and ggplot theme:
```{r message = F}
library(tidyverse)
library(lubridate)
library(raster)
library(oce) # for time series transformations
source("code/ggplot_theme.R")
**NOTE: This work flow uses many large files so most data is stored locally
and requires hard filepaths**
The local file path used for all large files is: `/Volumes/LACIE SHARE/
Stanford CH4/upch4 local/`.
Set the local head directory:
```{r}
loc <- "/Volumes/LACIE SHARE/Stanford CH4/upch4_local/"</pre>
Workflow
 1. Objective of Study
 2. Input Data
 + FLUXNET-CH4 Data
 + Gridded Data
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 + FLUXNET-CH4 Data Preparation
 - Averaging
 - Variable Selection
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 + Grid Extraction
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 - Computed (Rpot)
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 - N and S Deposition
 - SoilGrids
 - TerraClimate
```

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- + Merge Gridded Data
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- 6. Forward Feature Selection (FFS)
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- 7. Cross Validation
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    - RF and Final Predictors or Subsets
    - XGB and Final Predictors or Subsets
    - ANN and Final Predictors or Subsets
    - RNN and Final Predictors or Subsets
  - + Output Ensembles and Predictions
  - + Validation
    - Global Performance
    - Site-means
    - Monthly Seasonal Cycles
    - Monthly Anomalies
- 8. Variable Importances
  - + Variable Importance Rankings
  - + Variable Responses
  - + Partial Dependency Plots
  - + ShapR
- 9. Upscaled Model with Monte Carlo (MC)
  - + Forcing Data
    - Mapping
    - Member Product Choices
    - Evaluate Product Divergence
  - + Data Preparation
    - Extract Gridded Data for MC
    - Pre-Process FLUXNET-CH4 for MC
  - + MC Simulations
  - + MC ML Model Training
  - + MC ML Model Validation
- 10. Upscaling
  - + Prepare Member Forcing Data
  - + Run on Computing Cluster
    - Output Grids and Sums

- + Product Evaluation
  - Unweighted Wetland Fluxes (nmol)
  - Weighted Sums and Uncertainties (Tg)
  - Independent Validation
- 11. Data Representativeness
  - + Prepare Gridded Data
  - + Global Dissimilarity
  - + Tower Constituency
  - + Extrapolation Errors
    - MC ML Model Training Dissimilarity Only

[\*\*Link to Workflow Figure\*\*](https://drive.google.com/drive/folders/ 1jiFyqzoxxMpdtRLCwxCtzKpfILRNIW5K)

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## #### 1. Objective of Study

The goal of `FLUXNET-CH4 Upscaling` is to implement FLUXCOM-like ML approaches (e.g. [Jung et al. 2020](10.5194/bg-17-1343-2020)) to train a machine learning model using eddy covariance data that can predict wetland methane (CH4) fluxes globally. The predictions should be readily comparable to the Global Carbon Project (GCP) bottom-up process model ensembles that inform the Global Methane Budget ([Saunois et al. 2020](10.5194/ essd-12-1561-2020)). Wetland fluxes specifically, rather than methane fluxes from all terrestrial ecosystems, are the predictive goal of this study because 1) most eddy covariance data available are in wetlands, with limited coverage across the multitude of upland ecosystems, 2) methane fluxes are highest and most variable in wetlands, and 3) comparable bottomup process models predict wetland fluxes, then scale predictions to a global grid-cell using a prescribed (diagnostic runs) or model-derived (prognostic runs) wetland extent. In the last GCP Global Methane Budget ([Saunois et al. 2020](10.5194/essd-12-1561-2020)), diagnostic runs used the WAD2M product ([Zhang et al. in review](10.5194/essd-2020-262)). WAD2M includes coastal wetlands, however, we exclude coastal wetlands from the upscaling because they are salt-influenced and we do not have consistent salinity coverage, thus their inclusion is likely to bias flux estimates low even in non-coastal wetlands. The final model will be forced with available globally gridded data. Final product specifications are:

- Monthly time-step
- Historic reconstruction: ca. 2000 2018
- 0.25-degree grid cell resolution (as is WAD2M)
- Propagates training data uncertainties using Monte Carlo simulations
- Considers sensitivity to global forcing data product choices

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#### 2. Input Data

\*\*FLUXNET-CH4 Data\*\*

The eddy covariance data used in this study are publicly available as part of the FLUXNET-CH4 community product V1.0 at [fluxnet.org](https://fluxnet.org/data/fluxnet-ch4-community-product/). The FLUXNET-CH4 synthesis activity is introduced in [Knox et al. 2019](10.1175/BAMS-D-18-0268.1) along with a detailed description of the eddy covariance post-processing steps including methane flux (FCH4) uncertainties. The first full (V1.0) dataset release (FLUXNET-CH4, hereafter) is described for 81 sites and is used in a wetland seasonality analysis in [Delwiche et al. in review] (10.5194/essd-2020-307). \*\*Data for this CH4 Upscaling Project were downloaded from [fluxnet.org](www.fluxnet.org) on Feb 22, 2021. [Download Manifest](https://docs.google.com/spreadsheets/d/1--\_XyBqsyiMIdc6JXqhilOOYFLeaY-UTMFhhI4Sd5M/edit#qid=0)\*\*

## Links:

- [FLUXNET-CH4 Site (81) Metadata](https://docs.google.com/spreadsheets/d/1DN0huLs-vVM3g XcF1hBQrTpkKaGhzuWwaacfbe4iCo/edit#gid=1384338468)
- Permission was received via email on Feb 22, 2021, to use \*Tier 2 Data Policy\* sites in this study (SE-St1 and RU-Vrk; PI Thomas Friborg)
- FLUXNET-CH4 data were used both for methane fluxes (FCH4; target variable) and tower-measured bio-meteorological variables (e.g., LE, GPP, TA; predictors)
- + Link to [FLUXNET.org variable descriptions](https://fluxnet.org/
  data/fluxnet-ch4-community-product/data-variables/)

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\*\*Gridded Data (Predictors)\*\*

A summary of predictors (`Predictor Summary`) is available [here](https:// docs.google.com/spreadsheets/d/1DN0huLsvVM3g\_XcF1hBQrTpkKaGhzuWwaacfbe4iCo/edit#gid=0), as well as an appendix table with all individual predictors (`Appendix: All Predictors`). Candidate predictors were drawn from a mix of source classes (EC tower measurements, global models, computations from observed data, or remote sensing (e.g. NASA MODIS)), different information content groups (Spatialonly or Spatio-temporal), and at different temporal frequencies (Static, Yearly, Monthly, Weekly, or Half-hourly). Using this information, we assigned each candidate predictor to a class (Generic, Climate, Biometeorological, Land Cover, Soil and Relief, or Greenness) to evaluate predictive power of particular classes, given the likelihood for redundancy in useful predictors across the full predictor set. For MODIS predictors, we computed the mean seasonal cycle (`\_msc`), and yearly mean, min, max, and amplitude parameters, as in [Tramontana et al. (2016)](10.5194/ bg-13-4291-2016) and [Jung et al. (2020)](10.5194/bg-17-1343-2020). For derived TerraClimate soil moisture and actual evapotranspiration (aet) we computed the interannual range and annual seasonality. More information on preprocessing of predictors is provided in the manuscript.

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#### 3. FLUXNET-CH4 Pre-Processing

##### Local machine steps:

```
+ Create a copy of FLUXNET-CH4 data and name it `/fluxnet-ch4-data-
original`
 + Unzip all site flux files in `/fluxnet-ch4-data`
 + Reorganize into half-hourly (`/hh`) and `/daily` folders for easy
Look at one site of **daily means** data.
```{r}
# setwd(loc)
# files <- list.files("fluxnet-ch4-data/daily/")</pre>
# one_site <- read_csv(paste(loc, "fluxnet-ch4-data/daily/", files[1], sep</pre>
= ""))
# head(one_site)
No, but there is a quality flag `_QC`. `1`= data gap shorter than 2 months,
`3` = gap exceeds 2 months.
Issue: **There is also no uncertainty (`_UNC`) estimate on the downloaded
daily mean data.**
##### Clean up workspace:
```{r}
rm(one_site)
Look at one site of **half-hourly** data.
```{r}
setwd(loc)
files <- list.files("fluxnet-ch4-data/hh/")</pre>
one_site <- read_csv(paste(loc, "fluxnet-ch4-data/hh/", files[1], sep =
""))
head(one_site)
  - Half-hourly data has `FCH4` uncertainty columns `FCH4 F RANDUNC` and
`FCH4 F ANNOPTLM UNC`. Can be averaged over day or week.
  - Missing values are filled with `-9999`
##### Get FLUXNET-CH4 site names:
```{r}
site.names <- paste(substr(files, 5, 6), substr(files, 8, 10), sep = "")</pre>
Get all **half-hourly** flux data: (this will take a few minutes)
```{r echo = F, message = F, warning = F}
hh <- lapply(paste(loc, "fluxnet-ch4-data/hh/", files, sep = ""), read_csv)</pre>
names(hh) <- site.names</pre>
```

```
# str(hh)
##### Create a pristine replicate:
```{r}
hh2 <- hh
Look at all column names (including TIMESTAMP columns, to see date
and time format):
```{r}
names(hh[[1]])
head(hh[[1]]$TIMESTAMP_END)
##### Write and apply function to expand `TIMESTAMP END` into`Year`,
`Month`, `Week`, `Day`, and `DOY` variables to facilitate merging with
other data:
```{r}
expand date <- function(hh data) {
 hh_data <- hh_data %>%
 mutate(Year = as.numeric(substr(TIMESTAMP END, 1, 4)),
 Month = as.numeric(substr(TIMESTAMP_END, 5, 6)),
 Day = as.numeric(substr(TIMESTAMP_END, 7, 8)),
 Date = make date(Year, Month, Day),
 DOY = yday(Date),
 Week = ceiling(DOY/7),
 Week = ifelse(Week == 53, 52, Week)) %>%
 group_by(Year, DOY) %>%
 mutate(HH = 1:n()) %>%
 dplyr::select(Year, Month, Week, Day, HH, DOY, everything(), - Date, -
TIMESTAMP_START, -TIMESTAMP_END)
hh <- lapply(hh, expand_date)</pre>
hh[[2]] # check it worked
Check if gap-filled FCH4 `CH4_F_ANNOPTLM` has already been pre-filled
with observations `FCH4`, where available.
```{r}
hh[[1]]
FCH4 in row 7 in `hh[[1]]` (ID BRNpw) = 10.71, which **matches**
`CH4 F ANNOPTLM` = 1.071e+01.
**`CH4 F ANNOPTLM` has been pre-filled with observations.**
##### Write and apply function to create `imputed`, a flag variable where:
```

```
- `1` == `CH4_F_ANNOPTLM` was imputed
- `2` == `CH4_F_ANNOPTLM` was observed
```{r}
create_imputed <- function(hh_data) {</pre>
 hh data <- hh data %>%
 mutate(imputed = ifelse(FCH4 == -9999, 1, 0))
}
hh <- lapply(hh, create_imputed)</pre>
hh[[1]] check it worked
Create ID column:
```{r}
for (i in 1:length(hh)){
 hh[[i]] <- hh[[i]] %>%
    mutate(ID = site.names[i]) %>%
    dplyr::select(ID, everything())
head(hh[[1]])
##### Convert missing values (`-9999`) to `NA`:
```{r}
for (i in 1:length(hh)){
 hh[[i]][hh[[i]] == -9999] <- NA
head(hh[[1]])
Get `u` and `v` wind components:
Identify which sites lack WD:
```{r}
find_wd <- function(hh_data) {</pre>
  sum(names(hh data) == "WD") == 0
unlist(lapply(hh, find_wd))
sum(unname(unlist(lapply(hh, find_wd))))
OK, `CASCB` and `RUVrk` and missing `WD`.
Create dummy `WD` variables:
```

```
```{r}
hh$CASCB$WD <- NA
hh$RUVrk$WD <- NA
unlist(lapply(hh, find wd))
sum(unname(unlist(lapply(hh, find wd))))
Now compute wind direction components:
```{r}
compute_uv <- function(hh_data) {</pre>
  hh data <- hh data %>%
    mutate(U = -WS_F * sin(2 * pi * WD/360),
           V = -WS F * cos(2 * pi * WD/360))
}
hh <- lapply(hh, compute_uv)</pre>
head(hh[[1]]) # check it worked
##### Compute daily means of everything, including vector average WD and
speed, excluding precip. (sums)
Create function to compute daily means:
```{r echo = F}
compute daily1 <- function(hh data) {</pre>
 hh_data <- hh_data %>%
 group by(ID, Year, DOY) %>%
 summarize_all(list(~mean(., na.rm = T))) %>%
 mutate(WD_mean = (atan2(U, V) * 360/2/pi) + 180,
 WS mean = ((U^2 + V^2)^0.5)) %>%
 dplyr::select(ID, Year, Month, Week, Day, DOY, everything(), -HH)
}
Check run time for one site:
```{r warning = F}
system.time(
compute_daily1(hh[[1]])
) . .
Apply to all sites using lapply:
```{r warning = F}
```

```
daily <- lapply(hh, compute daily1)
Get sum of precip.
```{r warning = F, message = F}
sum precip <- function(hh data) {</pre>
  hh_data <- hh_data %>%
    group_by(ID, Year, DOY) %>%
    summarize(P_F_sum = sum(P_F))
}
daily_precip <- lapply(hh, sum_precip)</pre>
Rejoin precip. to main data frame:
```{r}
for (i in 1:length(daily)) {
daily[[i]] <- daily[[i]] %>%
 left_join(daily_precip[[i]], by = c("ID", "Year", "DOY"))
names(daily[[1]])
Output each "complete" daily site as a separate .csv file:
```{r}
for (i in 1:length(daily)) {
  daily[[i]] %>% write.csv(paste(loc, "fluxnet-ch4-data/daily_upch4/",
site.names[i], "_daily_upch4.csv", sep = ""),
                           row.names=FALSE)
}
#### 3. FLUXNET-CH4 Pre-Processing (cont.)
##### (re)Load Daily Data
```{r message = F, warning = F, echo = F}
files <- list.files(paste(loc, "fluxnet-ch4-data/daily_upch4/", sep = ""))</pre>
daily <- lapply(paste(loc,"fluxnet-ch4-data/daily_upch4/",files, sep = ""),</pre>
read_csv)
Bind rows
daily_flat <- daily %>% bind_rows() %>% as_tibble()
head(daily_flat)
Compute FCH4 uncertainty, and subset gap-filled (and best) predictors
Notes on variable selection:
```

```
+ Total FCH4 uncertainty `FCH4_F_UNC` is random `FCH4_F_RANDUNC` and gap-
filling `FCH4 F ANNOPTLM UNC` uncertainty summed in quadrature. I also drop
the `FCH4 F ANNOPTLM QC` Quality control flag because I implement a
stricter filtering criteria later of at least one observed flux per day.
 + The output of the daytime method `_DT` is used for gross primary
production (GPP) and ecosystem respiration (RECO). Although both methods
are subject to bias due to light-inhibition of leaf respiration, [Keenan et
al. 2019](https://doi.org/10.1038/s41559-019-0809-2) show that the biases
for the DT method only impact night-time respiration, but did not impact
apparent photosynthesis or daytime respiration.
 + LE and NEE were gap-filled according to [Knox et al. 2019](10.1175/
BAMS-D-18-0268.1) - same overall method to FCH4.
 + Other micro-meteorology (e.g., PPFD IN) was gap-filled using ERA
interim data (see [Delwiche et al.](10.5194/essd-2020-307)).
 + Only PPFD IN (not PPFD OUT) was selected as it can be approximated from
 + The shallowest available soil temperature is taken (`TS 1`)
```{r}
daily_subset <- daily_flat %>%
 mutate(FCH4_F_UNC = sqrt(FCH4_F_ANNOPTLM_UNC^2 + FCH4_F_RANDUNC^2)) %>%
 dplyr::select(ID, Year, Month, Week, Day, DOY, # descriptive data
         FCH4 = FCH4_F_ANNOPTLM, FCH4_F_UNC, imputed,
                                                        # methane fluxes
        NEE = NEE F ANNOPTLM, GPP = GPP DT, RECO = RECO DT, # ecosystem
C fluxes
        PPFD IN = PPFD IN F, SW IN = SW IN F, LW IN = LW IN F, NETRAD =
NETRAD F, # radiation
        LE = LE F ANNOPTLM, H = H F, # ecosystem energy fluxes
         TA = TA_F, PA = PA_F, RH = RH_F, VPD = VPD_F, P = P_F_sum, USTAR = PA_F
USTAR, WS = WS_mean, # meteorology
         TS = TS_1, SWC = SWC_F, WTD = WTD_F) # soil properties
head(daily_subset)
##### Append site metadata
Pulled from [FLUXNET-CH4 Site Metadata](https://docs.google.com/
spreadsheets/d/1DN0huLs-vVM3g XcF1hBQrTpkKaGhzuWwaacfbe4iCo/
edit#gid=1384338468)
metadata <- read_csv(paste(loc, "fluxnet-ch4-data/metadata/fluxnet-ch4-</pre>
site-metadata.csv", sep = ""))
daily subset meta <- metadata %>%
 mutate(ID = paste(substr(ID,1,2),substr(ID,4,6), sep="")) %>%
right_join(daily_subset, by = ("ID"))
##### Save subset, flattened, and metadata-appended daily FLUXNET-CH4 data
```{r}
write.csv(daily subset meta,
 paste(loc, "fluxnet-ch4-data/daily flat/daily subset meta.csv",
```

sep = ""),

```
row.names = F)
4. Gridded Data Pre-Processing
Potential Radiation (RPot)
Description
Get the mean seasonal cycle from Zutao Ouyang's 2001-2018 MATLAB output.
Original citation for `Rpot` is [Peltola et al. 2019](10.5194/
essd-11-1263-2019) where it was found to be a useful predictor of high
latitude wetland FCH4.
Define a function to take a number of years and return an index of months:
```{r}
create msc index <- function(years) {</pre>
  msc index <- list()</pre>
  for (i in 1:12){
      msc_index[[i]] \leftarrow seq(i, ((years-1)*12+i), by = 12)
  }
  msc_index
}
# create_msc_index(15) # test it out
Load in Rpot and create msc index for 19 years.
```{r}
Rpot <- brick(paste(loc, "grids/computed/Rpot.nc", sep = ""))</pre>
msc_index <- create_msc_index(19)</pre>
Get ` msc`:
```{r}
Rpot_msc <- list()</pre>
for (i in 1:12) {
 Rpot msc[[i]] <- calc(Rpot[[msc index[[i]]]], fun = mean)</pre>
}
Create raster stack:
```{r}
Rpot_msc <- stack(Rpot_msc)</pre>
Output `Rpot_msc.nc`:
```{r}
# writeRaster(Rpot_msc, paste(loc, "grids/computed/Rpot_msc.nc", sep = ""),
format="CDF", overwrite = T)
```

```
#### *MODIS*
**Description**
Zutao Ouyang (Stanford University) extracted MODIS pixels at the FLUXNET-
CH4 sites in April 2020, for 9 products:
 + Daytime Land Surface Temperature (`LSTD` from **MOD11A2**)
      - *(not used as it is a correlate of nighttime temp., and nighttime
is more applicable to soil conditions)*
 + Nighttime Land Surface Temperature (`LSTN` from **MOD11A2**)
 + Normalized Difference Vegetation Index (`NDVI` from **MOD09A1**)
 + Enhanced Vegetation Index (`EVI` from **MOD09A1**)
 + Leaf Area Index (`LAI` from **MCD15A2H**)
      - This is modeled, not directly measured.
 + Long Short Water Index (`LSWI` from **MOD09A1**)
 + Simple Ratio Water Index (`SRWI` from **MOD09A1**)
 + Normalized Difference Water Index (`NDWI` from **MOD09A1**)
 + Normalized Difference Snow Index (`NDSI` from **MOD11A2**)
**NOTE** Tables docmenting MODIS processing step effects on data are here
under [MODIS Processing](https://docs.google.com/spreadsheets/d/1DN0huLs-
vVM3g XcF1hBQrTpkKaGhzuWwaacfbe4iCo/edit#gid=1971246167) and QC figures are
output to `upch4_local/modis/modis_qc`.
##### Get file local file names/paths for 8-day extracted MODIS data:
```{r}
setwd(paste(loc, "/modis/modis-extracted", sep = ""))
files <- list.files()
Set MODIS product/file names (for gather values):
```{r}
modis.names <- c("LSTD", "EVI", "LAI", "LSWI", "NDSI", "NDVI", "NDWI",</pre>
"LSTN", "SRWI")
##### Read files:
```{r echo = F, warning = F, message = F}
```

modis <- lapply(paste(loc, "/modis/modis-extracted/", files, sep = ""),</pre>

read csv)

str(modis)

```{r}

head(modis[[1]])

names(modis) <- modis.names</pre>

Look at head for first file:

```
Files are not tidy (short and wide). There are **86** extracted sites
(extra sites than in FLUXNET-CH4 V1.0).
##### Use gather to make long and narrow, remove hyphen from ID:
```{r}
for (i in 1:length(files)) {
 modis[[i]] <- modis[[i]] %>%
 gather(key = ID, value = "modis.name", 2:87) %>%
 mutate(ID = paste(substr(ID, 1, 2), substr(ID, 4, 6), sep = "")) %>%
 as_tibble()
}
names(modis) <- modis.names</pre>
Check data are same length:
```{r}
str(modis)
1) `NDSI` is longer, 2) `LAI` is slightly shorter. Based on the dates,
`NDSI` is daily.
Will not be able to bind columns with different lengths
##### Remove `LAI` and `NDSI` then rejoin:
```{r}
modis.1.names <- c("LSTD", "EVI", "LSWI", "NDVI", "NDWI", "LSTN", "SRWI")</pre>
modis.1 <- modis[modis.1.names]</pre>
Select only first two columns (Date and ID) then data columns:
```{r}
modis.1 <- modis.1 %>%
  bind cols() %>%
  dplyr::select(1, 2, 3,6,9,12,15,18,21) %>% # subset only data columns
  as tibble()
names(modis.1) <- c("Date", "ID", modis.1.names)</pre>
##### Rejoin `LAI` using `Date`:
```{r}
modis.2 <- modis.1 %>% left_join(modis$LAI)
names(modis.2)[10] <- "LAI"</pre>
Rejoin `NDSI`:
```{r}
modis <- modis.2 %>% right_join(modis$NDSI)
```

```
names(modis)[11] <- "NDSI"</pre>
##### Convert `Date` into `Year` and `DOY`, correct `MARC` name (should be
`US-MAC`), remove `LSTD`:
```{r}
modis <- modis %>%
 mutate(Date = as.Date(Date, format = "%m/%d/%Y")) %>%
 mutate(ID = ifelse(ID == "MARC", "USMAC", ID),
 DOY = yday(Date),
 Date = as date(Date),
 Week = ceiling(DOY/7),
 Week = ifelse(Week == 53, 52, Week),
 Week = as.factor(Week),
 Year = as.integer(substr(Date, 1,4)),
 Month = as.numeric(substr(Date, 6, 7))) %>%
 dplyr::select(ID, Date, Year, Month, Week, DOY, NDSI, NDVI, EVI, LAI,
NDWI, SRWI, LSWI, LSTN)
Count `NAs` before despiking:
```{r}
modis nas <- modis %>%
 group_by(ID) %>%
  summarize all(list(~sum(is.na(.))))
modis nas
##### Despike outlier values for each product:
```{r}
modis_dsp <- modis %>%
 mutate(NDSI dsp = despike(NDSI, reference = 'trim', min = 0, max = 100,
replace = "NA"),
 NDVI dsp = despike(NDVI, reference = 'trim', min = -1, max = 1,
replace = "NA"),
 EVI_dsp = despike(EVI, reference = 'trim', min = -1, max = 1,
replace = "NA"),
 LAI dsp = despike(LAI, reference = 'trim', min = 0, max = 5,
replace = "NA"),
 NDWI_dsp = despike(NDWI, reference = 'trim', min = 0, max = 1,
replace = "NA"),
 SRWI_dsp = despike(SRWI, reference = 'trim', min = -1, max = 3,
replace = "NA"),
 LSWI_dsp = despike(LSWI, reference = 'trim', min = 0, max = 1,
replace = "NA"),
 LSTN_dsp = despike(LSTN, reference = 'trim', min = -60, max = 50,
replace = "NA")) %>%
dplyr::select(ID, Date, Year, Month, Week, DOY, NDSI, NDVI, EVI, LAI, NDWI,
SRWI, LSWI, LSTN,
 NDSI_dsp, NDVI_dsp, EVI_dsp, LAI_dsp, NDWI_dsp, SRWI_dsp,
LSWI_dsp, LSTN_dsp)
```

```
Count `NAs` after despiking:
```{r}
modis_dsp_nas <- modis_dsp %>%
 group by(ID) %>%
 summarize all(list(~sum(is.na(.))))
modis_dsp_nas
##### Calculate removed values during despiking and output to `upch4_local/
modis/modis qc/`
```{r}
modis dsp nas %>%
 mutate(NDSI = NDSI - NDSI_dsp,
 NDVI = NDVI - NDVI_dsp,
 EVI = EVI - EVI dsp,
 LAI = LAI - LAI_dsp,
 NDWI = NDWI - NDWI_dsp,
 SRWI = SRWI - SRWI_dsp,
 LSWI = LSWI - LSWI_dsp,
 LSTN = LSTN - LSTN_dsp) %>%
 dplyr::select(-ID) %>%
 summarize all(list(~sum(.))) %>%
 write.csv(paste(loc, "/modis/modis-qc/despiking na effects.csv", sep =
""))
. . .
Rename without `_dsp` suffix:
```{r}
modis_dsp <- modis_dsp %>%
 dplyr::select(ID, Date, Year, Month, Week, DOY,
                NDSI = NDSI_dsp,
                NDVI = NDVI dsp,
                EVI = EVI_dsp,
                LAI = LAI_dsp,
                NDWI = NDWI dsp,
                SRWI = SRWI dsp,
                LSWI = LSWI dsp,
                LSTN = LSTN dsp)
##### Calculate mean seasonal cycle (msc), then weekly means, then fill
weekly gaps with msc:
modis gapfilled <- modis dsp %>%
 group_by(ID, Month) %>% # this section computes monthly values averaged
over multiple years
 mutate(NDSI_msc = mean(NDSI, na.rm = TRUE),
         NDVI_msc = mean(NDVI, na.rm = TRUE),
```

```
EVI msc = mean(EVI, na.rm = TRUE),
         LAI_msc = mean(LAI, na.rm = TRUE),
         NDWI msc = mean(NDWI, na.rm = TRUE),
         SRWI_msc = mean(SRWI, na.rm = TRUE),
         LSWI_msc = mean(LSWI, na.rm = TRUE),
         LSTN msc = mean(LSTN, na.rm =TRUE)) %>%
 group by(ID, Year, Month, Week) %>% # this section compute weekly means
  summarize(NDSI = mean(NDSI, na.rm = TRUE),
            NDVI = mean(NDVI, na.rm = TRUE),
            EVI = mean(EVI, na.rm = TRUE),
            LAI = mean(LAI, na.rm = TRUE),
            NDWI = mean(NDWI, na.rm = TRUE),
            SRWI = mean(SRWI, na.rm = TRUE),
            LSWI = mean(LSWI, na.rm = TRUE),
            LSTN = mean(LSTN, na.rm = TRUE),
            NDSI msc = NDSI msc[1],
                                       # this section selects the monthly
(msc) value corresponding to the weekly data
            NDVI \ msc = NDVI \ msc[1],
            EVI_msc = EVI_msc[1],
            LAI msc = LAI msc[1],
            NDWI_msc = NDWI_msc[1],
            SRWI msc = SRWI msc[1],
            LSWI msc = LSWI msc[1],
            LSTN msc = LSTN msc[1]) %>%
 mutate(NDSI F = ifelse(is.na(NDSI), NDSI msc, NDSI), # this section fills
any missing weekly values with the msc (monthly average)
         NDVI_F = ifelse(is.na(NDVI), NDVI_msc, NDVI),
         EVI_F = ifelse(is.na(EVI), EVI_msc, EVI),
         LAI F = ifelse(is.na(LAI), LAI msc, LAI),
         NDWI_F = ifelse(is.na(NDWI), NDWI_msc, NDWI),
         SRWI_F = ifelse(is.na(SRWI), SRWI_msc, SRWI),
         LSWI_F = ifelse(is.na(LSWI), LSWI_msc, LSWI),
         LSTN_F = ifelse(is.na(LSTN), LSTN_msc, LSTN))
##### If snow is on the ground, set water indices to the 5% quantile
(frozen):
```{r}
modis_frozen <- modis_gapfilled %>%
 group by(ID,Year) %>%
 mutate(NDWI_msc = ifelse(NDSI_msc > 0, quantile(NDWI_msc, 0.05,
na.rm=TRUE), NDWI_msc),
 SRWI msc = ifelse(NDSI msc > 0, quantile(SRWI msc, 0.05,
na.rm=TRUE), SRWI msc),
 LSWI msc = ifelse(NDSI msc > 0, quantile(LSWI msc, 0.05,
na.rm=TRUE), LSWI_msc),
 NDWI F = ifelse(NDSI F > 0, quantile(NDWI F, 0.05, na.rm=TRUE),
NDWI F),
 SRWI F = ifelse(NDSI F > 0, quantile(SRWI F, 0.05, na.rm=TRUE),
SRWI F),
```

```
LSWI F = ifelse(NDSI F > 0, quantile(LSWI F, 0.05, na.rm=TRUE),
LSWI_F))
Compute mean, min, max, amplitude:
```{r message = F, warning = F}
modis_frozen <- modis_frozen %>%
  group_by(ID, Year) %>%
 mutate(NDSI_mean = mean(NDSI_F, na.rm=TRUE),
         NDVI_mean = mean(NDVI_F, na.rm=TRUE),
         EVI_mean = mean(EVI_F, na.rm=TRUE),
         LAI_mean = mean(LAI_F, na.rm=TRUE),
         NDWI mean = mean(NDWI F, na.rm=TRUE),
         SRWI_mean = mean(SRWI_F, na.rm=TRUE),
         LSWI_mean = mean(LSWI_F, na.rm=TRUE),
         LSTN_mean = mean(LSTN_F, na.rm=TRUE),
         NDSI_min = min(NDSI_F, na.rm=TRUE),
         NDVI_min = min(NDVI_F, na.rm=TRUE),
         EVI_min = min(EVI_F, na.rm=TRUE),
         LAI_min = min(LAI_F, na.rm=TRUE),
         NDWI min = min(NDWI F, na.rm=TRUE),
         SRWI_min = min(SRWI_F, na.rm=TRUE),
         LSWI min = min(LSWI F, na.rm=TRUE),
         LSTN_min = min(LSTN_F, na.rm=TRUE),
         NDSI_max = max(NDSI_F, na.rm=TRUE),
         NDVI max = max(NDVI F, na.rm=TRUE),
         EVI_max = max(EVI_F, na.rm=TRUE),
         LAI_max = max(LAI_F, na.rm=TRUE),
         NDWI_max = max(NDWI_F, na.rm=TRUE),
         SRWI_max = max(SRWI_F, na.rm=TRUE),
         LSWI_max = max(LSWI_F, na.rm=TRUE),
         LSTN_max = max(LSTN_F, na.rm=TRUE),
         NDSI_amp = NDSI_max-NDSI_min,
         NDVI_amp = NDVI_max-NDVI_min,
         EVI amp = EVI max-EVI min,
         LAI amp = LAI max-LAI min,
         NDWI_amp = NDWI_max-NDWI_min,
         SRWI_amp = SRWI_max-SRWI_min,
         LSWI_amp = LSWI_max-LSWI_min,
         LSTN_amp = LSTN_max-LSTN_min)
##### Get site IDs:
site.names <- modis frozen %>%
 ungroup() %>%
 mutate(ID = factor(ID)) %>%
 dplyr::select(ID) %>%
 pull() %>% unique()
```

```
##### Get variable names:
```{r}
names <- names(modis)[7:14]</pre>
names F <- paste(names, " F", sep = "")</pre>
names_msc <- paste(names, "_msc", sep = "")</pre>
Create modis qcqa figures:
```{r message = F, warning = F}
for (i in 1:length(names)){
  # visualize 1-45
  modis frozen %>%
    filter(ID %in% site.names[1:45]) %>%
    dplyr::select(Month, value_F = names_F[i], value_msc = names_msc[i])
%>%
    ggplot(aes(Month, value_F)) +
    geom_point(size = 1, alpha = 0.3) +
    geom line(aes(Month, value msc), col = 'purple', size = 2) +
    facet wrap(~ID, scales = 'free', ncol = 9) +
    my theme
  ggsave(paste(loc, "/modis/modis-qc/", names[i], "_1.pdf", sep = ""),
         width = 50, height = 30, units = c("cm"), dpi = 300)
  # visualize 46-86
  modis frozen %>%
    filter(ID %in% site.names[46:86]) %>%
    dplyr::select(Month, value_F = names_F[i], value_msc = names_msc[i])
%>%
    ggplot(aes(Month, value F)) +
    geom_point(size = 1, alpha = 0.3) +
    geom_line(aes(Month, value_msc), col = 'purple', size = 2) +
    facet_wrap(~ID, scales = 'free', ncol = 9) +
  ggsave(paste(loc, "/modis/modis-qc/", names[i], "_2.pdf", sep = ""),
         width = 50, height = 30, units = c("cm"), dpi = 300)
}
##### Output processed MODIS data:
write.csv(modis_frozen, paste(loc, "/modis/modis-processed/modis-
processed.csv", sep = ""),
          row.names = FALSE)
#### *Grid Extraction*
```

```
**Extract geospatial data at FLUXNET-CH4 sites and output a single
geospatial .csv.**
##### Load site metadata (ID, Latitude, Longitude), applicable to all
extractions:
```{r}
sites <- read_csv(paste(loc, "fluxnet-ch4-data/metadata/fluxnet-ch4-site-</pre>
metadata.csv", sep = "")) %>%
 mutate(ID = paste(substr(ID, 1, 2), substr(ID, 4, 6), sep = ""))
site.coords <- cbind(sites$Longitude, sites$Latitude) # (x, y)</pre>
site.num <- length(site.coords[,1])</pre>
Global Canopy Height
```{r}
raster.names <- list.files(paste(loc, "grids/global-canopy-height/", sep =
""), pattern = "tif$", full.names = FALSE)
stack <- stack(paste(loc, "grids/global-canopy-height/", raster.names, sep</pre>
= ""))
canopyht <- as_tibble(raster::extract(stack, site.coords))</pre>
canopyht <- cbind(sites, canopyht) %>%
  rename(canopyht = Simard Pinto 3DGlobalVeg JGR)
head(canopyht)
##### Computed (Rpot)
```{r}
stack <- stack(paste(loc, "grids/computed/Rpot_msc.nc", sep = ""))</pre>
Rpot <- as_tibble(raster::extract(stack, site.coords))</pre>
Rpot <- cbind(sites, Rpot) %>%
 gather(key = "Month", value = "Rpot", 15:26) %>%
 mutate(Month = str remove(Month, "X"))
head(Rpot)
Compound Topographic Index
```{r}
raster.names <- list.files(paste(loc, "grids/geomorpho90m/", sep = ""),</pre>
pattern = "tif$", full.names = FALSE)
stack <- stack(paste(loc, "grids/geomorpho90m/", raster.names, sep = ""))</pre>
cti <- as tibble(raster::extract(stack, site.coords))</pre>
cti <- cbind(sites, cti) %>%
  rename(cti = dtm cti merit.dem m 250m s0cm 2018 v1.0)
head(cti)
##### Earth Environment Texture, Land Cover and Topography
Texture (14 variables)
Land Cover (2 variables)
```

```
Topography (10 variables)
See: [Appendix: All Predictors](https://docs.google.com/spreadsheets/d/
1DN0huLs-vVM3g XcF1hBQrTpkKaGhzuWwaacfbe4iCo/edit#gid=0). Search for
`EarthEnv` under column `Gridded Product`.
###### First do texture:
```{r}
raster.names <- list.files(paste(loc, "grids/earthenv/texture/", sep = ""),</pre>
pattern = "tif$", full.names = FALSE)
stack <- stack(paste(loc, "grids/earthenv/texture/", raster.names, sep =</pre>
earthenv_texture <- as_tibble(raster::extract(stack, site.coords))</pre>
names(earthenv_texture) <-</pre>
c("cont", "corr", "cv", "diss", "entr", "even", "homo", "max", "rang", "shan", "simp", "std", "unif
Then land cover (commented code for first-run creates `HD` - human
development - as the sum of LC7 and LC9):
raster.names <- list.files(paste(loc, "grids/earthenv/cover/", sep = ""),
pattern = "tif$", full.names = FALSE)
stack <- stack(paste(loc, "grids/earthenv/cover/", raster.names, sep = ""))</pre>
create `HD` (human development)
HD <- stack[[1]] + stack[[2]]
writeRaster(HD, paste(loc, "/grids/earthenv/cover/HD.tiff", overwrite =
earthenv cover <- as tibble(raster::extract(stack, site.coords))</pre>
names(earthenv_cover) <- c("HD7", "HD9", "HD")</pre>
Then Topography:
```{r}
raster.names <- list.files(paste(loc, "grids/earthenv/topography/", sep =
""), pattern = "tif$", full.names = FALSE)
stack <- stack(paste(loc, "grids/earthenv/topography/", raster.names, sep =</pre>
earthenv_topography <- as_tibble(raster::extract(stack, site.coords))</pre>
names(earthenv topography) <-</pre>
c("east", "elev", "flat", "hollow", "north", "pcurv", "rough", "slope", "tcurv", "tpi")
###### Now combine:
earthenv <- cbind(sites, earthenv texture, earthenv cover,
earthenv topography)
###### Look at data histograms:
```

```
```{r}
earthenv %>%
 gather(key = "var", value = "value", 15:41) %>%
 ggplot(aes(value)) +
 geom histogram() +
facet_wrap(~var, scales = 'free')
`cont`, `diss`, and `var` have extreme outliers (xmax = 10e+09)
Replace extreme outliers with median
```{r}
earthenv <- earthenv %>%
  gather(key = "var", value = "value", 15:41) %>%
  group by(var) %>%
 mutate(value = ifelse(value > 10^9, median(value, na.rm = TRUE), value))
%>%
  ungroup() %>%
 spread(key = "var", value = "value")
Visualize again above to check they are corrected.
###### Also check for missing data:
```{r}
earthenv %>%
 gather(key = "var", value = "value", 15:41) %>%
 mutate(missing = is.na(value)) %>%
 group_by(ID) %>%
 summarize(total = n(),
 missing = sum(missing)) %>%
 arrange(desc(missing))
There are two sites (USDPW and USWPT) with missing values. Plot the
sites:
```{r}
plot(stack[[1]])
points(site.coords[sites$ID %in% c("USDPW", "USWPT"), ])
###### Fill with nearby site values.
USDPW from USLA1
USWPT from USOWC
```{r}
```

```
earthenv[earthenv$ID == "USDPW", c(is.na(earthenv[earthenv$ID == "USDPW",
]))] <- earthenv[earthenv$ID == "USLA1", c(is.na(earthenv[earthenv$ID ==
"USDPW",]))]
earthenv[earthenv$ID == "USWPT", c(is.na(earthenv[earthenv$ID == "USWPT",
]))] <- earthenv[earthenv$ID == "USOWC", c(is.na(earthenv[earthenv$ID ==
"USWPT",]))]
N and S Deposition
From [Lamarque et al. 2013](10.5194/acp-13-7997-2013):
 - accmip_nhx_acchist_2000.nc
 - accmip noy acchist 2000.nc
 - accmip_sox_acchist_2000.nc
```{r}
raster.names <- list.files(paste(loc, "grids/ns-deposition/", sep = ""),
pattern = "nc$", full.names = FALSE)
stack <- stack(paste(loc, "grids/ns-deposition/", raster.names, sep = ""))</pre>
plot(stack[[3]])
ns_depo <- as_tibble(raster::extract(stack, site.coords))</pre>
ns depo <- cbind(sites, ns depo) %>%
 rename(nx_depo = Dry.deposition.NH3.NH4,
         ny dep = Dry.deposition.NOy,
         s_dep = Dry.deposition.S02.S04)
head(ns_depo)
##### SoilGrids
##### TerraClimate
##### Fractional Vegetation Cover (VCF)
##### Wetland Extent (WAD2M)
```

WorldClim 2.0