

Solution NEON Part 2

K Duffy

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Sorry all, I haven't had time to finish this solution due to recording some future lectures for our class. I've push up an incomplete version, I'll revise and add more tomorrow.

```
library(tidyverse)
library(dplyr, quietly=T)
library(lubridate)
library(neonUtilities)
library(httr)
library(jsonlite)
library(downloader)
source('/Users/kdw223/Research/katharynduffy.github.io/neon_token_source.R')
```

1. Use the answers that you've provided above to select a *single* NEON site.

e.g. ONAQ

2. Use the answers that you've provided above to select **3 NEON data products** from either the TOS, TIS or ARS (AOP) collection methods. Summarize each product with its NEON identifier, along with a summary.

In this example solution I'm going to look at 3 phenological-related data products. This will vary based on your research and interests

Product 1, TOS Data: **DP1.10055.001**: Plant phenology observations: phenophase status and intensity of tagged plants. This data product contains the quality-controlled, native sampling resolution data from in-situ observations of plant leaf development and reproductive phenophases, at **D15.ONAQ**. Here I will focus on the phenophase intensity data, which is a measure of how prevalent that particular phenophase is in the sampled plants.

Product 2, TIS Data: **DP1.00033.001** Phenology images: RGB and IR images of the plant canopy taken from an automated camera on the tower top (note, there is another camera to capture understory imagery which is provided as a separate data product). Images are collected every 15 minutes and closely follow protocols of the Phenocam Network.

Product 3, RS Data: **DP3.30015.001** Ecosystem structure: If you're lazy like me, maybe you'd want to pull the abstract first and write something based on that:

```
# Request data using the GET function & the API call
req <- GET("http://data.neonscience.org/api/v0/products/DP3.30015.001")
req.content <- content(req, as="parsed")
req.content$data$productAbstract
```

```
## [1] "Forests store and sequester a considerable proportion of the terrestrial global carbon budget."
```

3. Using the NEON Utilities package or the API pull in those data along with metadata.

```
#TOS Phenology Data
sitesOfInterest <- c("ONAQ")
```

```

dpid <- as.character('DP1.10055.001') #phe data

pheDat <- loadByProduct(dpID="DP1.10055.001",
                         site = sitesOfInterest,
                         package = "expanded",
                         check.size = FALSE,
                         token=NEON_TOKEN)

```

Just to ‘mix things up’ I’ll use the API for my imagery:

```

#TIS Phenology Imagery
# Request data using the GET function & the API call
req <- GET("http://data.neonscience.org/api/v0/products/DP1.00033.001")
req

## Response [https://data.neonscience.org/api/v0/products/DP1.00033.001]
##   Date: 2020-08-28 23:18
##   Status: 200
##   Content-Type: application/json; charset=UTF-8
##   Size: 161 kB

# View requested data
req.content <- content(req, as="parsed")
names(req.content$data)

## [1] "productCodeLong"           "productCode"
## [3] "productCodePresentation"    "productName"
## [5] "productDescription"        "productStatus"
## [7] "productCategory"           "productHasExpanded"
## [9] "productScienceTeamAbbr"     "productScienceTeam"
## [11] "productPublicationFormatType" "productAbstract"
## [13] "productDesignDescription"    "productStudyDescription"
## [15] "productBasicDescription"     "productExpandedDescription"
## [17] "productSensor"              "productRemarks"
## [19] "themes"                     "changeLogs"
## [21] "specs"                      "keywords"
## [23] "siteCodes"

req.content$data$productAbstract

## [1] "Phenology is the study of reoccurring life cycle events that are driven by environmental factors

Interesting! I actually can't pull the NEON phenology imagery data from the NEON API. Good thing I checked the abstract. If I were a student (and not a former member of the PhenoCam lab) I'd probably go to the website, search for data etc etc, and then months later realize there's a slick phenocamAPI (yay APIs!!) that I can use.

I'll jump a few chapters ahead and use that data:

library(phenocamapi)
# obtaining midday_images for NEON ONAQ
ONAQ_middays <- get_midday_list('NEON.D15.ONAQ.DP1.00033')
# see the first few rows
head(ONAQ_middays)

## [1] "http://phenocam.sr.unh.edu/data/archive/NEON.D15.ONAQ.DP1.00033/2016/12/NEON.D15.ONAQ.DP1.00033"
## [2] "http://phenocam.sr.unh.edu/data/archive/NEON.D15.ONAQ.DP1.00033/2016/12/NEON.D15.ONAQ.DP1.00033"

```

```

## [3] "http://phenocam.sr.unh.edu/data/archive/NEON.D15.ONAQ.DP1.00033/2016/12/NEON.D15.ONAQ.DP1.00033"
## [4] "http://phenocam.sr.unh.edu/data/archive/NEON.D15.ONAQ.DP1.00033/2016/12/NEON.D15.ONAQ.DP1.00033"
## [5] "http://phenocam.sr.unh.edu/data/archive/NEON.D15.ONAQ.DP1.00033/2016/12/NEON.D15.ONAQ.DP1.00033"
## [6] "http://phenocam.sr.unh.edu/data/archive/NEON.D15.ONAQ.DP1.00033/2016/12/NEON.D15.ONAQ.DP1.00033"

```

Okay, now I have links to go grab those ‘mid-day’ images.

BUT...

The way I was calling data from the `neonUtilities` package doesn’t work anymore, so I could ask the `phenocamapi` package what it wants:

```

phenocams=get_phenos()
head(phenocams)

```

```

##          site      lat      lon elev active utc_offset date_first
## 1: aafcottawacfiaf14e 45.29210 -75.76640   90   TRUE       -5 2020-04-27
## 2: aafcottawacfiaf14w 45.29210 -75.76640   90   TRUE       -5 2020-05-01
## 3:         acadia 44.37694 -68.26083  158   TRUE       -5 2007-03-15
## 4: aguatibiaeast 33.62200 -116.86700 1086 FALSE      -8 2007-08-16
## 5: aguatibianorth 33.60222 -117.34368 1090 FALSE      -8 2003-10-01
## 6:        ahwahnee 37.74670 -119.58160 1199   TRUE      -8 2008-08-28
##          date_last infrared
## 1: 2020-08-24           N Elizabeth Pattey <elizabeth.pattey@canada.ca>
## 2: 2020-08-24           N Elizabeth Pattey <elizabeth.pattey@canada.ca>
## 3: 2020-08-27           N Dee Morse <dee_morse@nps.gov>
## 4: 2019-01-25           N Ann E Mebane <amebane@fs.fed.us>
## 5: 2006-10-25           N
## 6: 2020-08-27           N Dee Morse <dee_morse@nps.gov>
##          contact2
## 1: Luc Pelletier <luc.pelletier3@canada.ca>
## 2: Luc Pelletier <luc.pelletier3@canada.ca>
## 3:         John Gross <John_Gross@nps.gov>
## 4: Kristi Savig <KSavig@air-resource.com>
## 5:
## 6:         John Gross <John_Gross@nps.gov>
##          site_description site_type
## 1: AAFC Site - Ottawa (On) - CFIA - Field F14 -East Flux Tower      II
## 2: AAFC Site - Ottawa (On) - CFIA - Field F14 -West Flux Tower      II
## 3: Acadia National Park, McFarland Hill, near Bar Harbor, Maine     III
## 4:             Agua Tibia Wilderness, California      III
## 5:             Agua Tibia Wilderness, California      III
## 6: Ahwahnee Meadow, Yosemite National Park, California      III
##          group camera_description camera_orientation flux_data
## 1:      <NA> Campbell Scientific CCFC            NE    TRUE
## 2:      <NA> Campbell Scientific CCFC            WNW   TRUE
## 3: National Park Service                  unknown      NE   FALSE
## 4:          USFS                  unknown      SW   FALSE
## 5:          USFS                  unknown      NE   FALSE
## 6: National Park Service                  unknown       E   FALSE
##          flux_networks flux_sitenames
## 1:      <list[0]>      <NA>
## 2:      <list[1]>      <NA>
## 3:      <list[0]>
## 4:      <list[0]>
## 5:      <list[0]>

```

```

## 6:      <list[0]>
##                                         dominant_species primary_veg_type
## 1: Zea mays, Triticum aestivum, Brassica napus, Glycine max          AG
## 2: Zea mays, Triticum aestivum, Brassica napus, Glycine max          AG
## 3:                                         DB
## 4:                                         SH
## 5:                                         SH
## 6:                                         EN
##   secondary_veg_type site_meteorology MAT_site MAP_site MAT_daymet MAP_daymet
## 1:          AG        TRUE     6.4    943    6.30    952
## 2:          AG        TRUE     6.4    943    6.30    952
## 3:          EN       FALSE     NA     NA    7.05   1439
## 4:                      FALSE     NA     NA   15.75   483
## 5:                      FALSE     NA     NA   16.00   489
## 6:          GR       FALSE     NA     NA   12.25   871
##   MAT_worldclim MAP_worldclim koeppen_geiger ecoregion landcover_igbp
## 1:      6.0        863      Dfb      8      12
## 2:      6.0        863      Dfb      8      12
## 3:      6.5       1303      Dfb      8       5
## 4:     14.9        504      Csa     11       7
## 5:     13.8        729      Csa     11       7
## 6:     11.8       886      Csb      6       8
##   dataset_version1
## 1:          NA
## 2:          NA
## 3:          NA
## 4:          NA
## 5:          NA
## 6:          NA
##
## 1: Camera funded by Agriculture and Agri-Food Canada (AAFC) Project J-001735 - Commercial inhibitor
## 2: Cameras funded by Agriculture and Agri-Food Canada (AAFC) Project J-001735 - Commercial inhibitor
## 3:
## 4:
## 5:
## 6:
##   modified flux_networks_name flux_networks_url
## 1: 2020-05-04T10:46:30.065790-04:00           <NA>           <NA>
## 2: 2020-05-04T10:46:32.523976-04:00           OTHER
## 3: 2016-11-01T15:42:15.016778-04:00           <NA>           <NA>
## 4: 2016-11-01T15:42:15.086984-04:00           <NA>           <NA>
## 5: 2016-11-01T15:42:15.095277-04:00           <NA>           <NA>
## 6: 2016-11-01T15:42:15.111916-04:00           <NA>           <NA>
##   flux_networks_description
## 1:           <NA>
## 2: Other/Unaffiliated
## 3:           <NA>
## 4:           <NA>
## 5:           <NA>
## 6:           <NA>

```

Looking at the dataframe and searching for ONAQ I see that I have to give this full name to pull the imagery:
`NEON.D15.ONAQ.DP1.00033`

```

# open a temporary directory
tmp_dir <- tempdir()

# download a subset.
download_midday_images(site = 'NEON.D15.ONAQ.DP1.00033', # which site
                       y = 2019, # which year(s)
                       months = 1:12, # which month(s)
                       days = 15, # which days on month(s)
                       download_dir = tmp_dir) # where on your computer

## | 
## [1] "/var/folders/5n/vxrdb72140b43_jhsmhf514pqgq1n/T//RtmpB1cEwV"
# list of downloaded files
ONAQ_middays_path <- dir(tmp_dir, pattern = 'NEON.D15.ONAQ.*', full.names = TRUE)

head(ONAQ_middays_path)

## [1] "/var/folders/5n/vxrdb72140b43_jhsmhf514pqgq1n/T//RtmpB1cEwV/NEON.D15.ONAQ.DP1.00033_2019_01_15"
## [2] "/var/folders/5n/vxrdb72140b43_jhsmhf514pqgq1n/T//RtmpB1cEwV/NEON.D15.ONAQ.DP1.00033_2019_02_15"
## [3] "/var/folders/5n/vxrdb72140b43_jhsmhf514pqgq1n/T//RtmpB1cEwV/NEON.D15.ONAQ.DP1.00033_2019_03_15"
## [4] "/var/folders/5n/vxrdb72140b43_jhsmhf514pqgq1n/T//RtmpB1cEwV/NEON.D15.ONAQ.DP1.00033_2019_04_15"
## [5] "/var/folders/5n/vxrdb72140b43_jhsmhf514pqgq1n/T//RtmpB1cEwV/NEON.D15.ONAQ.DP1.00033_2019_05_15"
## [6] "/var/folders/5n/vxrdb72140b43_jhsmhf514pqgq1n/T//RtmpB1cEwV/NEON.D15.ONAQ.DP1.00033_2019_06_15"

```

Great, the data is there.

4. Organize your data into data.frames and produce summaries for each of your data:

```

#NEON sends the data as a nested list, so I need to undo that
# unlist all data frames
list2env(pheDat ,.GlobalEnv)
summary(phe_perindividualperyear)
summary(phe_statusintensity)

```

5. Filter and format your data based on metadata and quality flags:

```

#remove duplicate records
phe_statusintensity <- select(phe_statusintensity, -uid)
phe_statusintensity <- distinct(phe_statusintensity)
#Format dates (native format is 'factor' silly R)
phe_statusintensity$date <- as.Date(phe_statusintensity$date, "%Y-%m-%d")
phe_statusintensity$editedDate <- as.Date(phe_statusintensity$editedDate, "%Y-%m-%d")
phe_statusintensity$year <- substr(phe_statusintensity$date, 1, 4)
phe_statusintensity$monthDay <- format(phe_statusintensity$date, format="%m-%d")

```

Now I want to remove NA values so I can see what's really going on:

```

phe_statusintensity=phe_statusintensity%>%
  filter(!is.na(phe_statusintensity$phenophaseIntensity))

```

Based on looking at the status intensity data and metadata I'm going to just stick to the `status.intensity:leaves` metrics here.

6. Create minimum of 1 plot per data type (minimum of 3 plots total). These will vary based on that data that you've chosen.

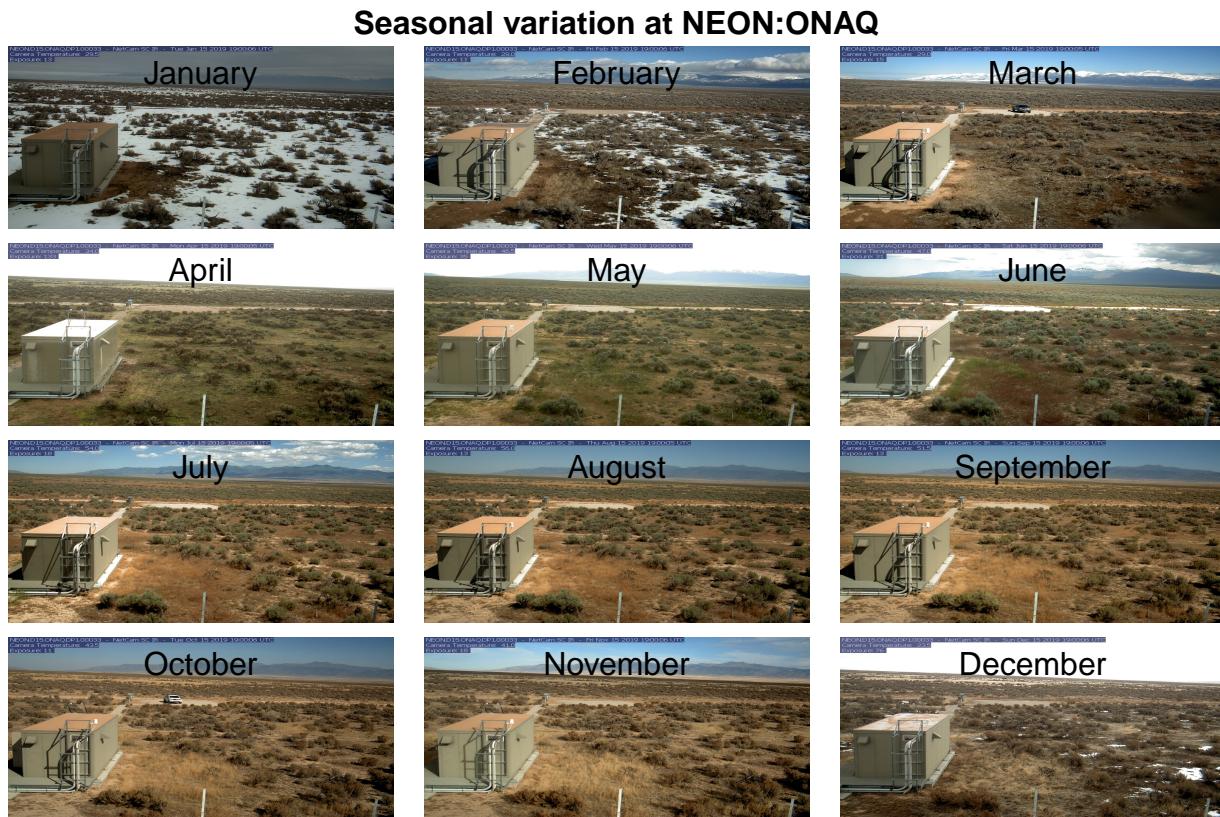
Second, let's look at the PhenoCam imagery over a year:

```

library(jpeg)
n <- length(ONAQ_middays_path)
par(mar= c(0,0,0,0), mfrw=c(4,3), oma=c(0,0,3,0))

for(i in 1:n){
  img <- readJPEG(ONAQ_middays_path[i])
  plot(0:1,0:1, type='n', axes= FALSE, xlab= '', ylab = '')
  rasterImage(img, 0, 0, 1, 1)
  mtext(month.name[i], line = -2)
}
mtext('Seasonal variation at NEON:ONAQ', font = 2, outer = TRUE)

```



7. What is the temporal frequency of observations in the data you decided was of interest? How do the data align to answer a central question? What challenges did you run into when investigating these data? How will you address these challenges and document your code? *One to two paragraphs*

Pulling the abstracts really helped me with this. The TOS phenology data is measured every

The PhenoCam imagery is taken every 15 minutes. Think about just how much that is!

The AOP flights are ideally every year, but that can be hard to coordinate (we'll talk about that in Chapter 7).