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#Table of Contents-----
# BH-01 starts on line 47
# BL-02 starts on line 192
# BH-02 start on line ...
#Set working directory to ensure R can find the files we wish to import.
setwd("C:/Users/aalda/Desktop/All plots 2018")
#Installing and Loading all packages-----
#install.packages("RStoolbox")
#install.packages("rasterVis")
#install.packages('raster')
#install.packages('gdalUtils')
#install.packages('tidyverse')
#install.packages('rgr')
#install.packages('uavRst')
#install.packages('rgdal') -Alex
library('gdalUtils')
library('RStoolbox')
library('rasterVis')
library('raster')
library('ggplot2')
library('rgr')
library('tidyverse')
library('rgdal') #alex wrote this.
#library('uavRst') #may not have been used.
# Load Data ------
                  # We will create a character vector list of raster files using the list.files() function in the directory
named "All plots 2018". This list will be used to generate a Rasterstack.
files <- list.files()</pre>
files
                  #See the list of all files in the directory named "All plots 2018".
dbf.files <- files[grep(".tiff", files, fixed=T)]  #Greates a file that is list of names only having .tif extensions. Grep function finds ".tiff" pattern in the created "files" and fixed=T means pattern is a text string.
for(i in dbf.files) { assign(unlist(strsplit(i, "[.]"))[1], raster(i)) }
BH1 RGB<-stack("BH-01 RGB modified.tif") # Import multi-band raster data, using the stack() function.
BH1_IR<-raster('BH-01 IR_modified.tif') # Import and create a Rasterlayer file using the raster function.
BL2_RGB<-stack("BL-02 RGB.tif")
BL2 IR<-raster('BL-02 IR.tif')
# BH-01 -----
#align extent
BH1_IR_proj<-projectRaster(BH1_IR, BH1_RGB) #Alex: Project the data of a Raster object to a new RAster object with another
projection (crs). projectRaster(from, to).
#Shapefile
plot(BH1_IR_proj)
                                   #press-clear broom stick in plots tab, if you get a error that margins are too large.
Alex+ani: line 58-Error-plot.new has not been called yet, occured becuase plot name has not been called.
polygon<-shapefile("Polygons.shp")  #Alex: File format for storing geospatial data in polygon.shp.
plot(polygon, add=TRUE)
                                   #Alex: This adds another raster ontop of another. This draws the boundary of the two
exclosures over the [BH1_IR_proj] image
BH1_shp_ug<-subset(polygon, PlotID=='BH-01 UG') #Ungrazed #Alex: Returns (selected variables) subsets of vectors, matirces or
dataframes which meet conditions. Subset(object to be subsetted, logical expression indicating elements or rows to keep).
BH1 shp g<-subset(polygon, PlotID=='BH-01 G') #Grazed
# Mask and clip rasters to polygon ------
#Ungrazed
BH1 RGB mask<-mask(BH1 RGB, BH1 shp ug)
BH1_IR_mask<-mask(BH1_IR_proj, BH1_shp_ug)
BH1RGB_crop<- crop(BH1_RGB_mask, BH1_shp_ug)
plot(BH1RGB crop)
                           #Alex: four images produced. (#1-image)
ex<-extent(BH1RGB_crop)
BH1IR_crop<- crop(BH1_IR_mask, ex)</pre>
plot(BH1IR crop)
                            #Alex: one image produced.(#2-image)
# Stack and Brick IR and RGB -----
BH1 stack<-stack(BH1RGB crop, BH1IR crop)
nlayers(BH1_stack)
BH1_stack<-writeRaster(BH1_stack, filename="C:/Users/aalda/Desktop/All plots 2018/BH1_Stack.tif", format="GTiff", overwrite=TRUE)
# Calculate NDVI -------
BH1\_ndvi\_ungrazed <- ((BH1\_stack[[5]]-BH1\_stack[[1]]) / (BH1\_stack[[5]]+BH1\_stack[[1]])) \\
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#Alex: one image produced.(#3-image)

plot(BH1\_ndvi\_ungrazed)

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# Do it all again for BH1 Grazed ------
# Mask and clip rasters to polygon -------
BH1 RGB mask<-mask(BH1 RGB, BH1 shp g)
BH1 IR mask<-mask(BH1 IR proj, BH1 shp g)
BH1RGB_crop<- crop(BH1_RGB_mask, BH1_shp_g)</pre>
                           #Alex: four images produced. (#5-image)
plot(BH1RGB crop)
ex<-extent(BH1RGB crop)
BH1IR_crop<- crop(BH1_IR_mask, ex)
plot(BH1IR crop)
                           #Alex: one image produced.(#6-image)
# Stack and Brick IR and RGB ------
BH1 stack<-stack(BH1RGB crop, BH1IR crop)
nlayers (BH1_stack)
                            #Alex: there are 5 layers.
# Calculate NDVI -----
BH1_ndvi_grazed<-((BH1_stack[[5]]-BH1_stack[[1]])/(BH1_stack[[5]]+BH1_stack[[1]]))
plot(BH1_ndvi_grazed)
                                #Alex: one image produced. (#7-image)
BL2 IR proj<-projectRaster(BL2_IR, BL2_RGB) #Alex: copied and pasted here.
BL2_shp_g<-subset(polygon, PlotID=='BL-02 G')#Grazed -Alex copied and pasted here.
BL2_IR_mask<-mask(BL2_IR_proj, BL2_shp_g) #Alex: first run line 181 for BL2_shp_g and line 185 for BL2_IR_proj.
EL12RGB_crop<- crop(BL2_RGB_mask, BL2_shp_g) #Alex: first run line 229 (BL2_RGB_mask<-mask(BL2_RGB_shp_g). Problem: Error in file(fn, "rb"): cannot open the connection. Solution: run line 72 and line 16 (library (raster)).
plot(BL2RGB crop)
                                                #Alex: four images in the shape of parallelgrams.(#8 image)
ex <- extent (BL2RGB crop)
BL2IR_crop<- crop(BL2_IR_mask, ex)
plot(BL2IR crop)
                                               #Alex: one image produce in the shape of parallelgram. (#9-image)
# Stack and Brick IR and RGB -----
BL2 stack<-stack(BL2RGB crop, BL2IR crop)
nlayers (BL2 stack)
BL2\_ndvi\_grazed < -((BL2\_stack[[5]] - BL2\_stack[[1]]) / (BL2\_stack[[5]] + BL2\_stack[[1]])) \\
plot(BL2_ndvi_grazed)
                                                 #Alex: one image produced in the shape of a parallelgram. (#10 image)
hist(BL2_ndvi_grazed)
                                                 #Alex: histogram produced. (#11 image)
#compare to ungrazed
BL2\_ndvi\_ungrazed <-((BL2\_stack[[5]]-BL2\_stack[[5]]+BL2\_stack[[5]]+BL2\_stack[[1]])) \\ \#alex:copied/pasted here.
BL2_ndvi_ungrazed<-as.data.frame(BL2_ndvi_ungrazed) #Alex: First run line 217 for BL2_ndvi_ungrazed which is (
BL2_ndvi_ungrazed<-((BL2_stack[[5]]-BL2_stack[[1]])/(BL2_stack[[5]]+BL2_stack[[1]]))
BL2_ndvi_grazed<-as.data.frame(BL2_ndvi_grazed)
BH_01_g<-tibble(
  Value=BL2_ndvi_grazed$layer,
Treatment="Grazed"
BH_01_ug<-tibble(
  Value=BL2_ndvi_ungrazed$layer,
Treatment="Ungrazed"
{\tt BH\_01 < -rbind(BH\_01\_g, BH\_01\_ug)}
ggplot(data=BH_01, aes(x=Treatment, y=Value))+
  geom_violin(scale='area')
                                                  #Alex: two images produced.(#12 image)
hist(BH1_ndvi_grazed)
                                                  #Alex: histogram produced. (#13 image)
#compare to ungrazed
BH1_ndvi_ungrazed<-as.data.frame(BH1_ndvi_ungrazed)</pre>
BH1_ndvi_grazed<-as.data.frame(BH1_ndvi_grazed)
BH_01_g<-tibble(
  Value=BH1_ndvi_grazed$layer,
Treatment="Grazed"
BH_01_ug<-tibble(
  Value=BH1_ndvi_ungrazed$layer,
Treatment="Ungrazed"
 \begin{array}{ll} {\tt BH\_01<-rbind(BH\_01\_g,\ BH\_01\_ug)} \\ {\tt ggplot(data=BH\_01,\ aes(x=Treatment,\ y=Value))+} \end{array} 
  geom_violin(scale='area')
                                                      #Alex: two images produced. (#14 image)
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#Alex: histogram produced. (#4-image)

hist(BH1 ndvi ungrazed)

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#align extent
BL2 IR proj<-projectRaster(BL2 IR, BL2 RGB) #Alex: two images produced. (#15 image)
#Shapefile
plot(BL2 IR proj)
                                               #Alex created this line.
polygon<-shapefile("Polygons.shp")</pre>
plot(polygon, add=TRUE)
BL2_shp_ug<-subset(polygon, PlotID=='BL-02 UG') #Ungrazed
BL2_shp_g<-subset(polygon, PlotID=='BL-02 G') #Grazed
# Wrong: BL2_shp_ug<-subset(polygon, PlotID=='BH-01 UG') #Ungrazed
                                                                             #Alex: warming messages.
#wrong: BL2_shp_g<-subset(polygon, PlotID=='BH-01 G')#Grazed</pre>
# Mask and clip rasters to polygon ------
#Ungrazed
BL2_RGB_mask<-mask(BL2_RGB, BL2_shp_ug)
BL2_IR_mask<-mask(BL2_IR_proj, BL2_shp_ug)
                                                    #Alex: first run line 191 for BL2_shp_ug. Problem: Error in .local(x, y, ...) :
BL2RGB_crop<- crop(BL2_RGB_mask, BL2_shp_ug)
extents do not overlap
plot(BL2RGB crop)
                                     #Alex: four images produced in shape of parallelgrams. (#16 image)
ex<-extent(BL2RGB crop)
BL2IR_crop<- crop(BL2_IR_mask, ex)</pre>
plot(BL2IR_crop)
                                       #Alex: (check) one image not produced. (#17 image)
# Stack and Brick IR and RGB ------
BL2_stack<-stack(BL2RGB_crop, BL2IR_crop)</pre>
nlayers (BL2 stack)
                                                 #Alex: 5 layers.
BL2_ndvi_ungrazed<-((BL2_stack[[5]]-BL2_stack[[1]])/(BL2_stack[[5]]+BL2_stack[[1]])) plot(BL2_ndvi_ungrazed) #Alex: (check) one image produced. (#18 image
                                            #Alex: (check) one image produced. (#18 image)
hist(BL2 ndvi ungrazed)
                                            #Alex: (check) histogram produced. (#19 image)
# Do it all again for BL2 Grazed ------
# Mask and clip rasters to polygon ------
#Grazed
BL2_RGB_mask<-mask(BL2_RGB, BL2_shp_g)
BL2_IR_mask<-mask(BL2_IR_proj, BL2_shp_g)
                                                  #Ales wrote this line.
#pasted here.
BL2RGB_crop<- crop(BL2_RGB_mask, BL2_shp_g)  #Alex: first run line 229 (BL2_RGB_mask<-mask(BL2_RGB, BL2_shp_g). Problem: Error in file(fn, "rb"): cannot open the connection. Solution: run line 72 and line 16 (library (raster)). plot(BL2RGB_crop)  #Alex: four images in the shape of parallelgrams.(#8 image)
ex<-extent(BL2RGB_crop)
BL2IR_crop<- crop(BL2_IR_mask, ex)
plot(BL2IR_crop)
                                                  #Alex: one image produce in the shape of parallelgram. (#9-image)
# Stack and Brick IR and RGB ------
BL2_stack<-stack(BL2RGB_crop, BL2IR_crop)
nlayers (BL2_stack)
# Calculate NDVI ------
\verb|BL2_ndvi_grazed<-((BL2\_stack[[5]]-BL2\_stack[[1]])/(BL2\_stack[[5]]+BL2\_stack[[1]]))|
plot(BL2_ndvi_grazed)
hist(BL2_ndvi_grazed)
                                                     #Alex: one image produced in the shape of a parallelgram.(#10 image)
                                                     #Alex: histogram produced. (#11 image)
#compare to ungrazed
BL2_ndvi_ungrazed<-((BL2_stack[[5]]-BL2_stack[[1]])/(BL2_stack[[5]]+BL2_stack[[1]])) #alex:copied/pasted here. BL2_ndvi_ungrazed<-as.data.frame(BL2_ndvi_ungrazed) #Alex: First run line 217 for BL2_ndvi_ungrazed which is (BL2_ndvi_ungrazed<-((BL2_stack[[5]]-BL2_stack[[1]])/(BL2_stack[[5]]+BL2_stack[[1]])) .
BL2_ndvi_grazed<-as.data.frame(BL2_ndvi_grazed)</pre>
#alex wrote this
BL 02 q<-tibble(
  Value= BL2 ndvi ungrazed$layer,
  Treatment="Grazed"
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```
BL_02_ug<-tibble(
    Value=BL2_ndvi_ungrazed$layer,
    Treatment= "Ungrazed"
)

BL_02<-rbind(BL_02_g, BL_02_ug)
ggplot(data= BL_02, aes (x= Treatment, y=Value))+
    geom_violin(scale='area')
hist(BL2_ndvi_grazed)  # Error in hist.default(BL2_ndvi_grazed) : 'x' must be numeric. Image#</pre>
```