

Ayata-et-al-2018

The objective of this section is to predict SOC across an area of interest based on terrain parameters and machine learning. We use 18 soil profile descriptions and a set of environmental information to predict the spatial variability of SOC across a water limited environment of Northeast Mexico, including its associated uncertainty.

data preparation: estimating SOC stocks 0-30cm depth

```
dat <- read.csv("horizon.csv")
```

```
sites <- read.csv("site.csv")
```

```
library(aqp)
```

```
## This is aqp 1.16
```

```
depths(dat) <- ID ~ top + bottom
```

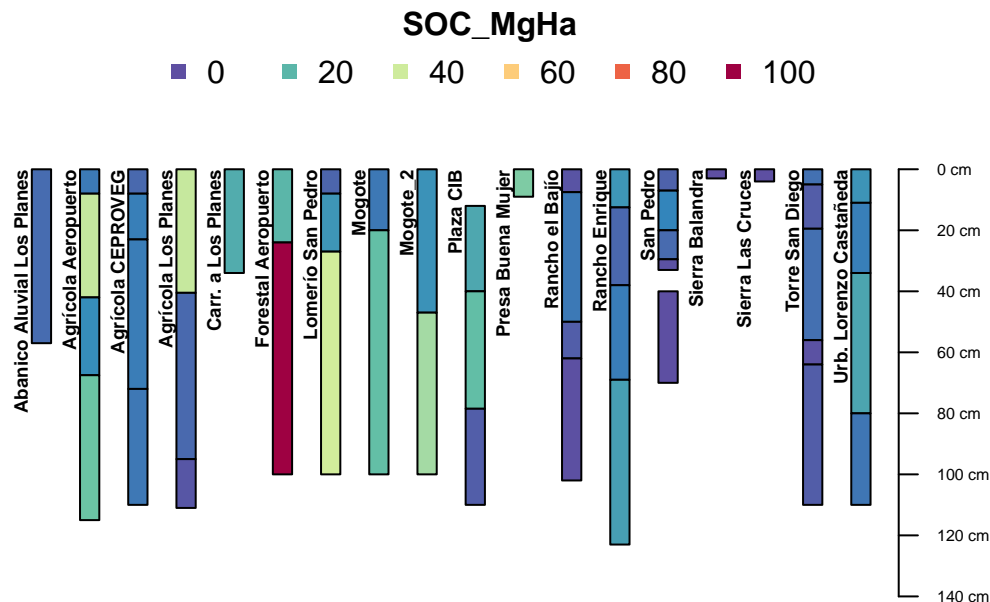
```
## Warning: converting IDs from factor to character
```

```
dataqp <- dat
```

```
#VISUALIZE SOC DATA
```

```
plot(dataqp, color='SOC_MgHa')
```

```
## unable to guess column containing horizon designations
```



```
site(dat) <- sites
```

```
coordinates(dat) <- ~ X + Y
```

```
library(GSIF)
```

```

## GSIF version 0.5-4 (2017-04-25)
## URL: http://gsif.r-forge.r-project.org/
try(OCS <- mpspline(dat, 'SOC_MgHa', d = t(c(0,30))))

## Fitting mass preserving splines per profile...
##
|
|
| 0%

## Spline not fitted to profile: Abanico Aluvial Los Planes
##
|
|====| 6%
|
|=====| 11%
|
|=====| 17%
|
|=====| 22%

## Spline not fitted to profile: Carr. a Los Planes
##
|
|=====| 28%
|
|=====| 33%
|
|=====| 39%
|
|=====| 44%
|
|=====| 50%
|
|=====| 56%

## Spline not fitted to profile: Presa Buena Mujer
##
|
|=====| 61%
|
|=====| 67%
|
|=====| 72%
|
|=====| 78%

## Spline not fitted to profile: Sierra Balandra
##
|
|=====| 83%

## Spline not fitted to profile: Sierra Las Cruces

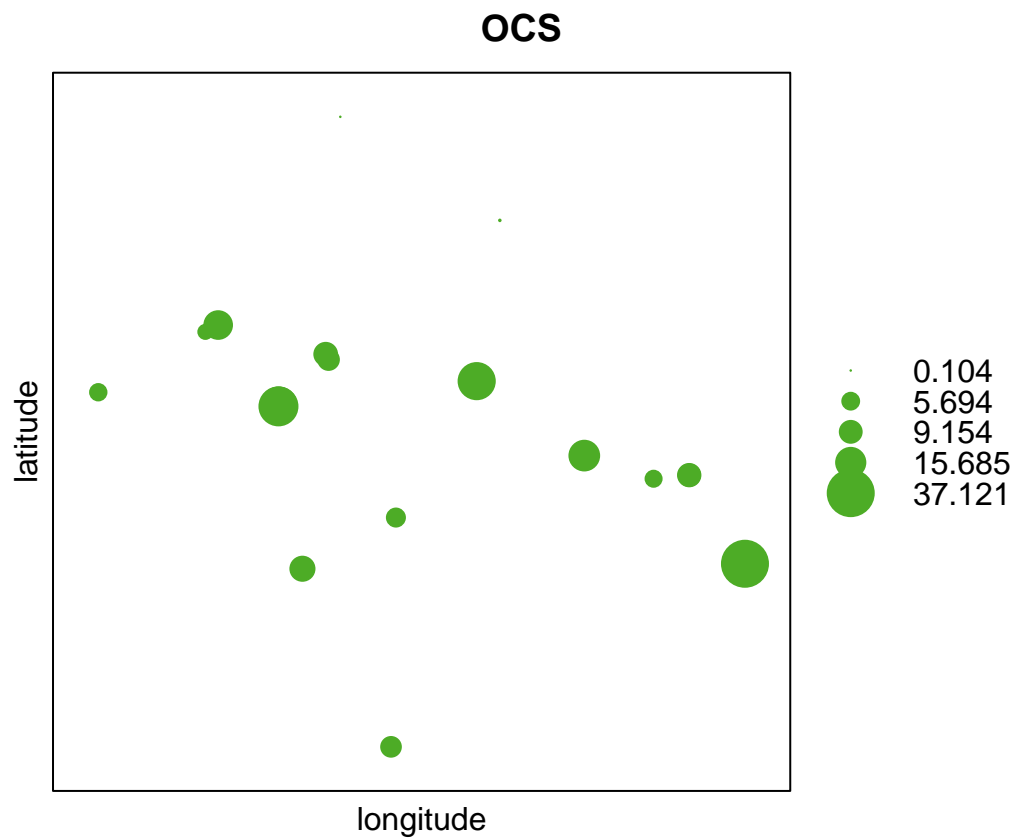
```

```
##
|=====| 89%
|=====| 94%
|=====| 100%
```

```
dat <- data.frame(id = dat@site$ID,
                  Y = dat@sp@coords[,2],
                  X = dat@sp@coords[,1],
                  OCS = OCS$var.std[,1])
```

```
#write.csv(dat, "mexico_data.csv")
```

```
datstp <- dat
coordinates(datstp) <- ~ Y + X
#SPATIAL LOCATIONS
library(sp)
bubble(datstp['OCS'], xlab='longitude', ylab='latitude')
```



prepare covariates: harmonize all available prediction factors

generate dummy variables for those categorical prediction factors

```
dummyRaster <-function(rast){
  rast <-as.factor(rast)
  result <-list()
  for(i in 1:length(levels(rast)[[1]][[1]])){
    result[[i]] <-rast==levels(rast)[[1]][[1]][i]
    names(result[[i]]) <-paste0(names(rast),
    levels(rast)[[1]][[1]][i])}
  return(stack(result))
}

library (raster)

#SELECT THE COLUMN NUMBERS OF INTEREST
#lis all tif files
#lis1 the separated maps (i.e., landforms)
#lis2 continuos maps (i.e., prec)
#lis3 categorical maps (i.e., soil type)
(lis <- list.files(pattern='tif$'))
(lis1 <- lis[-c(5, 9, 10, 11, 12, 13, 35, 41)])
(lis2 <- lis[c(35)])
(lis3 <- lis[c(9, 10, 13, 41)])
#AREA OF INTEREST
aoi <- raster("Area de estudio.tif")
#TOPOGRAPHIC TERRAIN PARAMETERS DERIVED ON SAGA GIS
dem <- stack('dem15/terrain/terrain.tif')
dem[is.na(dem)==TRUE]<- -9999
dem[is.infinite(dem)==TRUE]<- 9999
names(dem) <- c('dem','hillshade','curvature','convergenceIndex','flowAccumulation','wetnessIndex','l

dum <- stack()

for (i in 1:length(lis1)){
  r <- raster (lis1[i])
  r <- projectRaster (r, aoi)
  r <- crop(r, aoi)
  r[is.na(r)==FALSE,] <- 1
  r[is.na(r)==TRUE,] <- 10
  #r <- mask (r, aoi)
  dum <- stack(dum, r)
  print(paste0(i, names(r), ' done!'))
}

cont <- stack()

for (i in 1:length(lis2)){
  r <- raster (lis2[i])
  r <- projectRaster (r, aoi)
  r <- crop(r, aoi)
```

```

    #r <- mask (r, aoi)

    cont <- stack(cont, r)

    print(paste0(i, names(r), ' done!'))
  }

cat <- stack()

for (i in 1:length(lis3)){
  r <- raster (lis3[i])
  r[is.na(r)==TRUE,] <- -9999
  r <- projectRaster (r, aoi, method='ngb')
  r <- crop(r, aoi)
  #r <- mask (r, aoi)
  r <- dummyRaster(r)
  cat <- stack(cat, r)
  print(paste0(i, names(r), ' done!'))
}

cat$Edafología_Serie_II4[is.na(cat$Edafología_Serie_II4)==TRUE] <- 2
cat$Edafología_Serie_II6[is.na(cat$Edafología_Serie_II6)==TRUE] <- 2

COVS <- stack(dum, cont, cat)
COVS <- COVS[[-7]]
COVS[is.infinite(COVS)==TRUE] <- -9999
COVS[is.na(COVS)==TRUE] <- -9999
library(RStoolbox)
COVS <- scale(COVS)

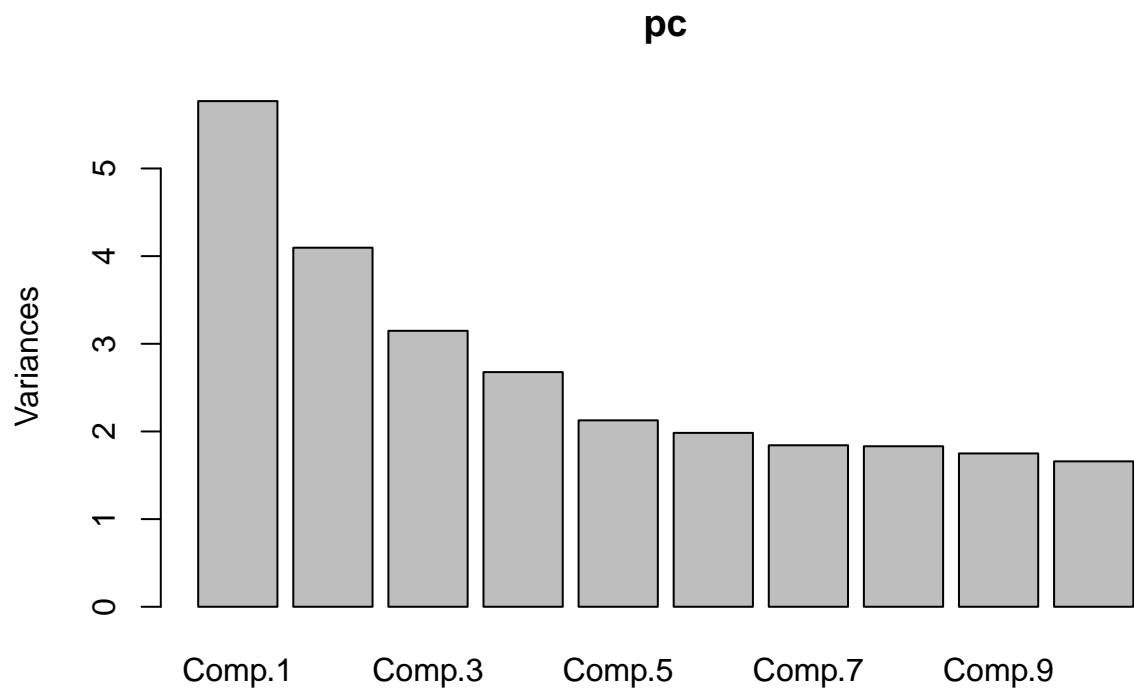
```

prepare predictors for PCA, include the terrain parameters to the covariate space and generate a regression matrix (couple with points of soil profiles)

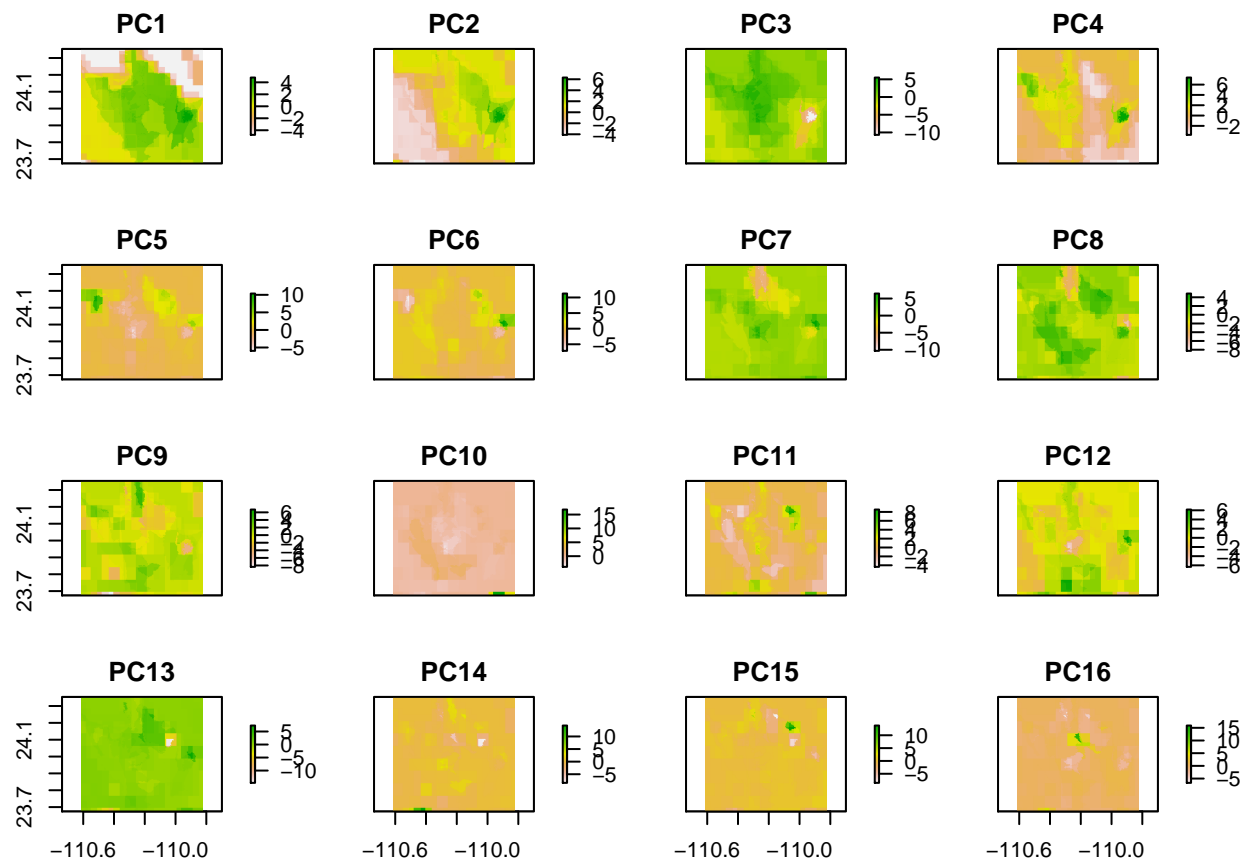
```

COVSpca <- rasterPCA(COVS, maskCheck=FALSE)
pc <- COVSpca$model
#PLOT PCA INERTIA
plot(pc)

```



```
#PLOT PCA maps
plot(stack(COVSpca$map))
```



```
COVSpca <- stack(COVSpca$map)[[1:3]]
```

```
x <- stack(resample( dem, COVSpca), COVSpca)

training <- cbind(data.frame(extract(x, datsp)), OCS = dat$OCS, Y = dat$X, X = dat$Y)

#summary(training)
#CHECK CORRELATED PREDICTORS
round(cor(training), 2)
```

```
##          dem hillshade curvature convergenceIndex
## dem          1.00      0.32      0.16           0.15
## hillshade     0.32      1.00      0.00          -0.08
## curvature      0.16      0.00      1.00           0.75
## convergenceIndex 0.15     -0.08      0.75           1.00
## flowAccumulation -0.29     -0.24      0.01           0.14
## wetnessIndex    -0.68     -0.25     -0.04           0.10
## lsFactor        -0.48     -0.06     -0.05           0.07
## slope           0.68      0.50     -0.21          -0.33
## aspect          0.12     -0.73     -0.01           0.01
## PC1             0.00      0.06     -0.09           0.25
## PC2             0.06      0.23      0.12           0.20
## PC3             0.14     -0.24     -0.14          -0.35
## OCS             0.07      0.02     -0.25           0.06
## Y              -0.23      0.21      0.10          -0.36
## X               0.27      0.29      0.16           0.31
##          flowAccumulation wetnessIndex lsFactor slope aspect  PC1
## dem          -0.29          -0.68    -0.48  0.68  0.12  0.00
## hillshade     -0.24          -0.25    -0.06  0.50 -0.73  0.06
## curvature       0.01          -0.04    -0.05 -0.21 -0.01 -0.09
## convergenceIndex 0.14           0.10     0.07 -0.33  0.01  0.25
## flowAccumulation 1.00           0.65     0.79 -0.26  0.36  0.19
## wetnessIndex     0.65           1.00     0.91 -0.75  0.14  0.25
## lsFactor         0.79           0.91     1.00 -0.45  0.20  0.25
## slope           -0.26          -0.75    -0.45  1.00 -0.05 -0.18
## aspect          0.36           0.14     0.20 -0.05  1.00 -0.10
## PC1             0.19           0.25     0.25 -0.18 -0.10  1.00
## PC2             0.18           0.04     0.20  0.12 -0.09  0.80
## PC3            -0.05          -0.24    -0.22  0.21  0.40 -0.74
## OCS            -0.24          -0.32    -0.46  0.06 -0.38  0.48
## Y              0.10          -0.13     0.03  0.32 -0.09 -0.46
## X              0.08          -0.04     0.12  0.20 -0.11  0.75
##          PC2  PC3  OCS  Y  X
## dem          0.06  0.14  0.07 -0.23  0.27
## hillshade     0.23 -0.24  0.02  0.21  0.29
## curvature      0.12 -0.14 -0.25  0.10  0.16
## convergenceIndex 0.20 -0.35  0.06 -0.36  0.31
## flowAccumulation 0.18 -0.05 -0.24  0.10  0.08
## wetnessIndex    0.04 -0.24 -0.32 -0.13 -0.04
## lsFactor        0.20 -0.22 -0.46  0.03  0.12
## slope           0.12  0.21  0.06  0.32  0.20
## aspect        -0.09  0.40 -0.38 -0.09 -0.11
## PC1            0.80 -0.74  0.48 -0.46  0.75
## PC2            1.00 -0.73  0.23 -0.04  0.89
## PC3           -0.73  1.00 -0.38  0.37 -0.76
## OCS            0.23 -0.38  1.00 -0.31  0.23
```

```
## Y          -0.04  0.37 -0.31  1.00 -0.30
## X          0.89 -0.76  0.23 -0.30  1.00
```

define color pallete for maps, remove non assigned values and mask the prediction space to the area of interest

```
jet.colors <- colorRampPalette(c("#00007F", "blue", "#007FFF", "cyan",
                                "#7FFF7F", "yellow", "#FF7F00", "red", "#7F0000"))

x <- stack(resample(COVSpca, dem), dem)
x[is.na(x)==TRUE]<- -9999
x[is.infinite(x)==TRUE]<- 9999

aoi <- resample(aoi, x, method='ngb')
x <- mask(x, aoi)
```

run 1 predictive model with all the 18 points

```
library(caret)

s <- stack()
m <- list()
r2 <- numeric()
rmse <- numeric()
#REPEATED CROSS-VALIDATION
control <- rfeControl(functions=rfFuncs, method="repeatedcv",          number=2, repeats=5)

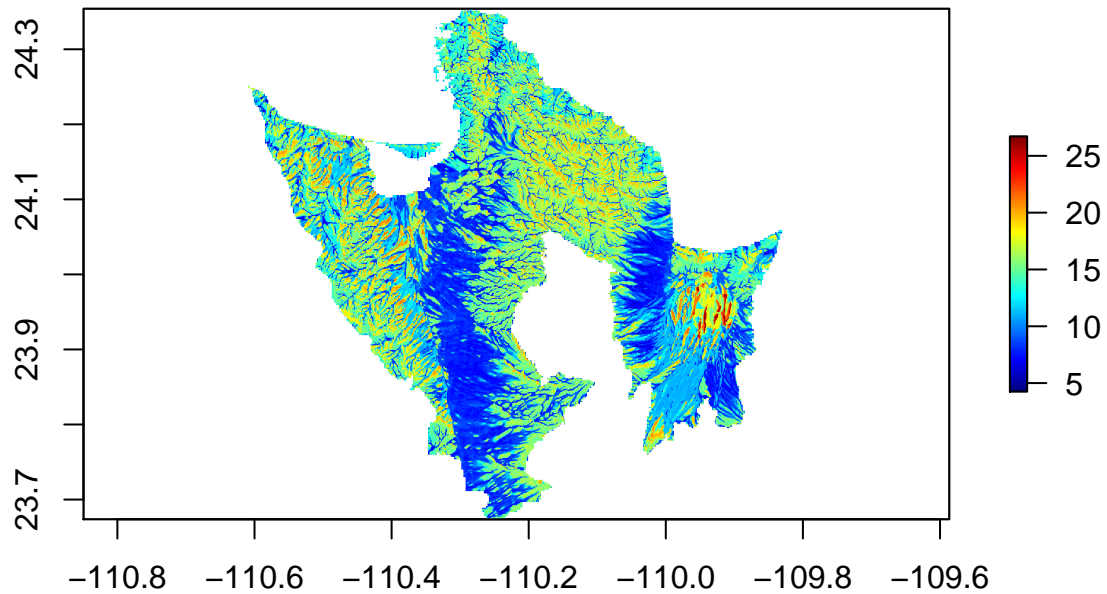
#10 MODELS FOR TESTING
for (i in 1:10){
  #RFE recursive feature elimination based on RANDOM FORESTS
  rfProfile <- rfe(training[,1:12], training[,13], sizes=c(1:12),          rfeControl=control)

  #BEST FIT
  m[[i]] <- rfProfile
  rmse[i] <- max(m[[i]]$results[2])
  r2[i] <- max(m[[i]]$results[3])

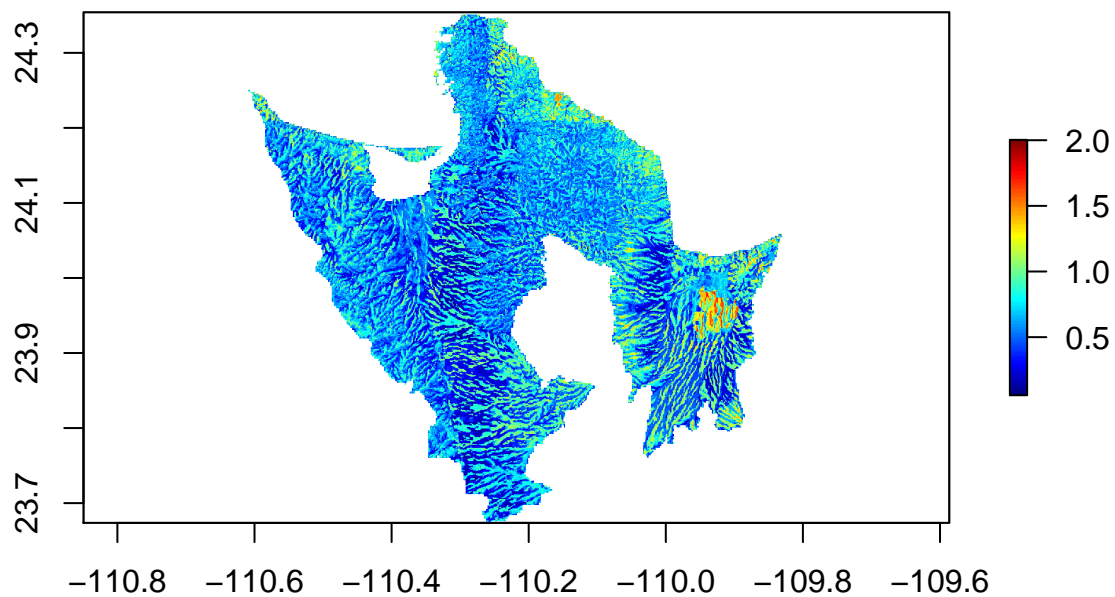
  print(rfProfile)
  predictors(rfProfile)
  predRFE <- predict(x, rfProfile)
  #plot(predRFE, col=jet.colors(100))
  s <- stack(s, predRFE)
  names(s)[[i]] <- paste0('model-', i)
}
```


plot the predicted maps and the variance map

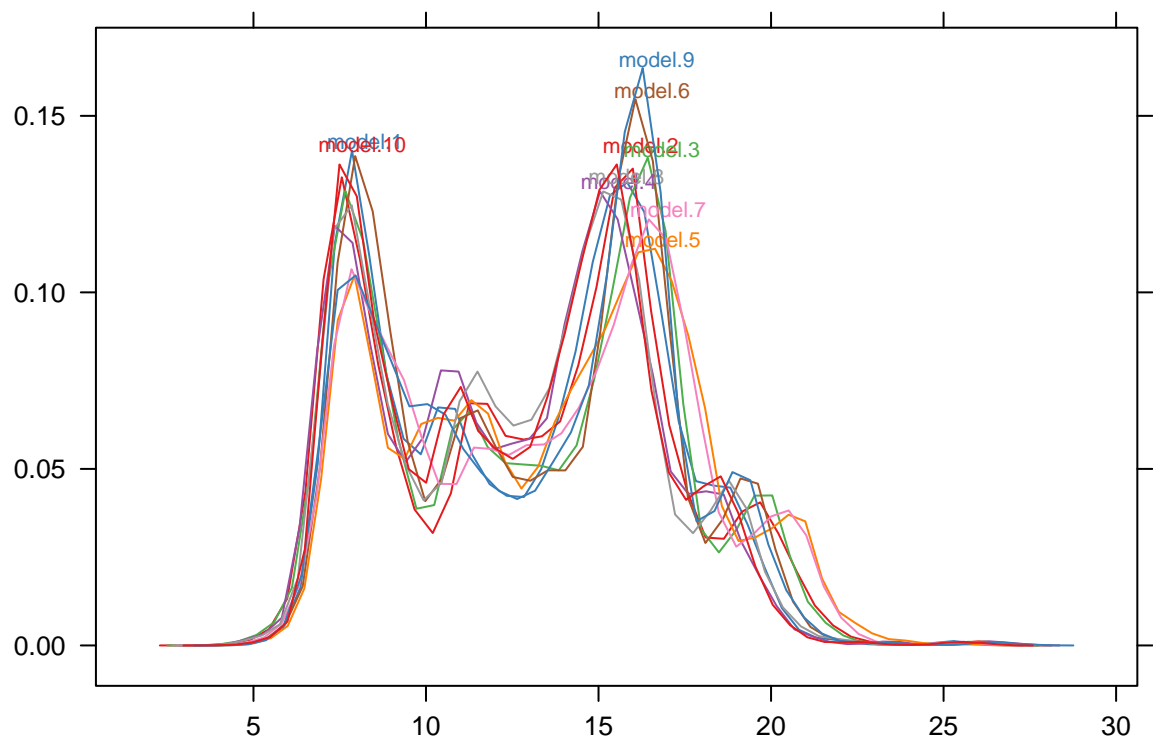
```
#MEAN PREDICTION  
plot(calc(s, mean), col=jet.colors(100))
```



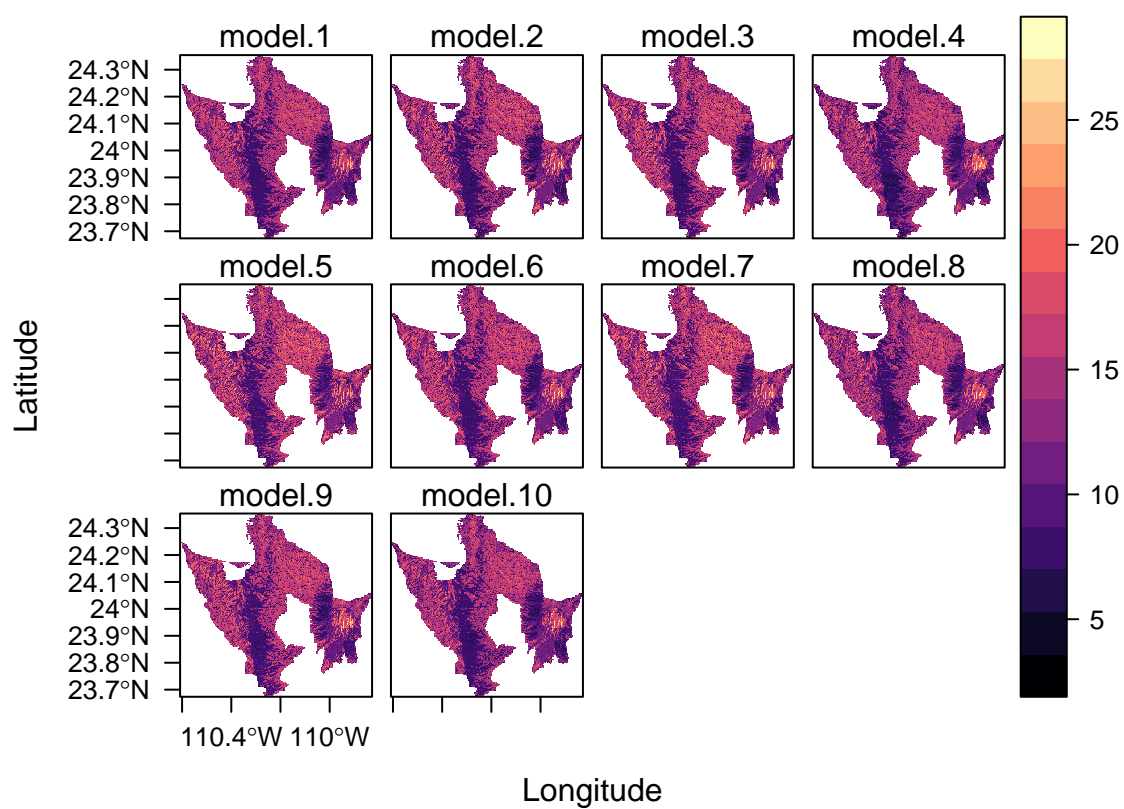
```
#PREDICTION VARIANCE (UNCERTAINTY)  
plot(calc(s, sd), col=jet.colors(100))
```



```
#UNCERTAINTY  
rasterVis::densityplot(s)
```



```
#ALL PREDICTIONS
rasterVis::levelplot(s)
```



```
#writeRaster(s, file='SOCpredictions.tif')
```

accuracy numbers

```
#EXPLAINED VARIANCE
```

```
summary(r2)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
## 0.2235  0.2804  0.3300  0.3121  0.3475  0.3535
```

```
#RMSE
```

```
summary(rmse)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##  9.756   9.977  10.300  10.350  10.520  11.510
```

```
#sum pixes and calculate the total SOC stocks for all the area
```

```
#cellStats(calc(s, mean), sum)
```

```
#and the uncertainty
```

```
#cellStats(calc(s, sd), sum)
```

around 30% of explained variance with a mean error of 9.9 Mg.Ha.

end of exercise