

# Q1: Are lake and terrestrial primary productivity coherent?

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This document organizes for openness and reproducibility analyses of the temporal coherence of interannual variation in lake primary productivity with terrestrial primary productivity in the landscape surrounding the lake.

## Data import

Data produced in ‘ms1\_prep.Rmd’ are loaded.

```
load("/Users/jonathanwalter/Box Sync/NSF EAGER Synchrony/Data/RData files/ms1_analysis_inprogress1_v108")

any(sapply(analysislakes$lakedata, function(x){any(is.infinite(x))}))

## [1] FALSE

any(sapply(analysislakes$lakedata, function(x){any(is.na(x))}))

## [1] FALSE

which(sapply(analysislakes$lakedata, function(x){any(is.na(x))}))

## named integer(0)

analysislakes$lakeinfo[which(sapply(analysislakes$lakedata, function(x){any(is.na(x))})),]

## [1] lagoslakeid      gnis_name      nhd_lat
## [4] nhd_long           lake_area_ha   lake_perim_meters
## [7] nhd_ftype          nhd_fcode      hu4_zoneid
## [10] hu12_zoneid        state_zoneid    elevation_m
## [13] start              end
## <0 rows> (or 0-length row.names)

# image(accndvi)
# points(lakepts.prj[which(sapply(analysislakes$lakedata, function(x){any(is.na(x))})),])

dbuff[which(sapply(analysislakes$lakedata, function(x){any(is.na(x))}))]

## numeric(0)

analysislakes$lakeinfo<-analysislakes$lakeinfo[!sapply(analysislakes$lakedata, function(x){any(is.na(x))}),]
analysislakes$lakedata<-analysislakes$lakedata[!sapply(analysislakes$lakedata, function(x){any(is.na(x))}),]

analysislakes$lakeinfo$tslength<-analysislakes$lakeinfo$end-analysislakes$lakeinfo$start+1
# analysislakes$lakedata<-analysislakes$lakedata[!analysislakes$lakeinfo$tslength < 20]
# analysislakes$lakeinfo<-analysislakes$lakeinfo[!analysislakes$lakeinfo$tslength < 20,]

source("~/GitHub/AquaTerrSynch/AnalysisCode/bandtest_coh.R")

tsranges<-rbind(c(2,4),c(4,Inf),c(2,Inf))
```

```

coh.chlaXaccndvi<-NULL
coh.chlaXmaxndvi<-NULL

for(lind in 1:length(analysislakes$lakedata)){
  lakedat.ii<-cleandat(analysislakes$lakedata[[lind]], as.numeric(colnames(analysislakes$lakedata[[lind]])))
  chlaXaccndvi<-coh(lakedat.ii[1,], lakedat.ii[2,], as.numeric(colnames(analysislakes$lakedata[[lind]])))
    norm="powall", sigmethod="fast", nrand=10000)
  chlaXmaxndvi<-coh(lakedat.ii[1,], lakedat.ii[3,], as.numeric(colnames(analysislakes$lakedata[[lind]])))
    norm="powall", sigmethod="fast", nrand=10000)
  for(rind in 1:nrow(tsranges)){
    chlaXaccndvi<-bandtest.coh(chlaXaccndvi, tsranges[rind,])
    chlaXmaxndvi<-bandtest.coh(chlaXmaxndvi, tsranges[rind,])
  }
  coh.chlaXaccndvi<-rbind(coh.chlaXaccndvi, c(t(as.matrix(chlaXaccndvi$bandp[,3:5]))))
  coh.chlaXmaxndvi<-rbind(coh.chlaXmaxndvi, c(t(as.matrix(chlaXmaxndvi$bandp[,3:5]))))
}

coh.chlaXaccndvi<-as.data.frame(coh.chlaXaccndvi)
coh.chlaXmaxndvi<-as.data.frame(coh.chlaXmaxndvi)

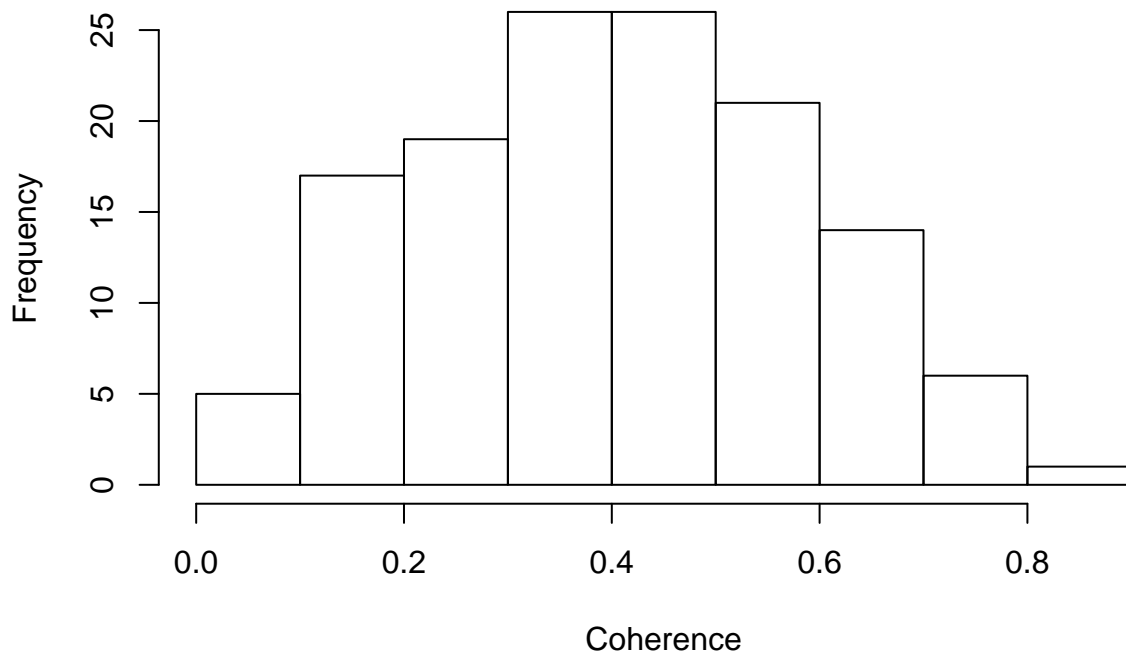
colnames(coh.chlaXaccndvi)<-paste0("accndvi",c("p.ts1","phi.ts1","coh.ts1","p.ts2","phi.ts2","coh.ts2",
colnames(coh.chlaXmaxndvi)<-paste0("maxndvi",c("p.ts1","phi.ts1","coh.ts1","p.ts2","phi.ts2","coh.ts2",

coh.chlaXaccndvi$lagoslakeid<-analysislakes$lakeinfo$lagoslakeid
coh.chlaXmaxndvi$lagoslakeid<-analysislakes$lakeinfo$lagoslakeid

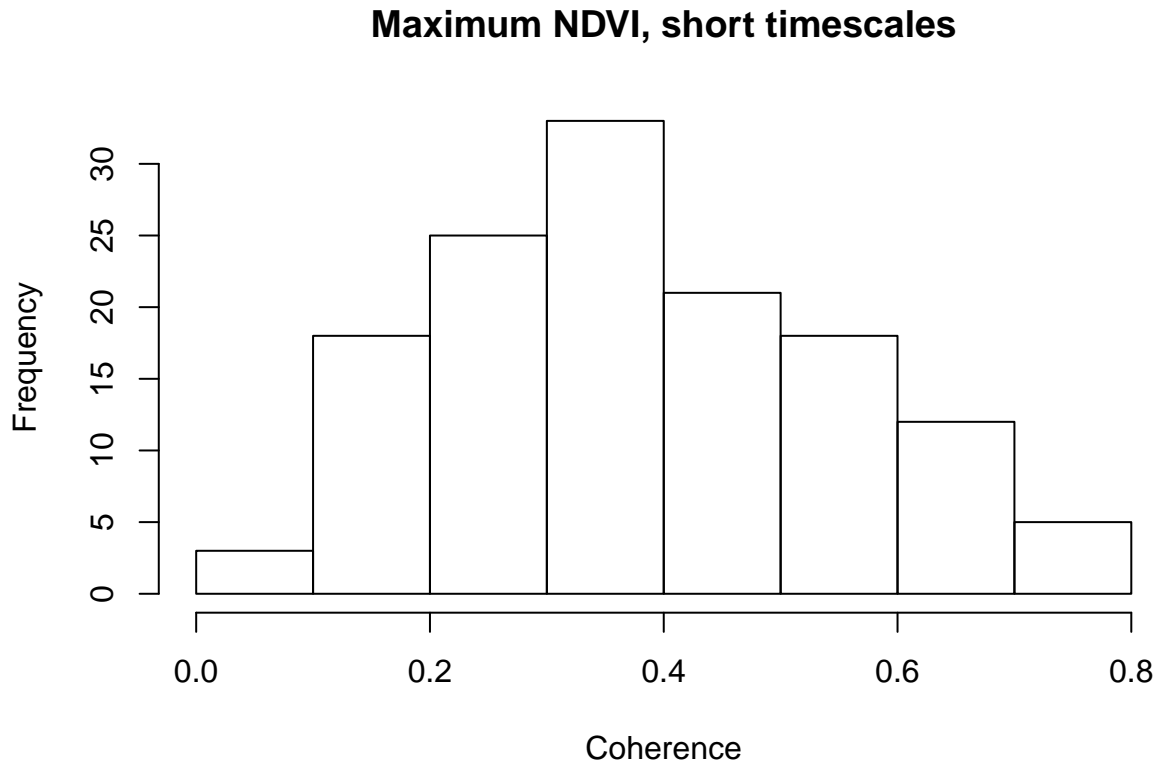
#short timescales
hist(coh.chlaXaccndvi$accndvicoh.ts1, main="Accumulated NDVI, short timescales", xlab="Coherence", ylab="Frequency", col="white", border="black")

```

## Accumulated NDVI, short timescales



```
hist(coh.chlaXmaxndvi$maxndvicoh.ts1, main="Maximum NDVI, short timescales", xlab="Coherence", ylab="Fr
```



```
quantile(coh.chlaXaccndvi$accndvicoh.ts1)
```

```
##          0%          25%          50%          75%         100%
## 0.03540956 0.26015941 0.40373548 0.52492077 0.81625251
```

```
quantile(coh.chlaXmaxndvi$maxndvicoh.ts1)
```

```
##          0%          25%          50%          75%         100%
## 0.04514692 0.24996954 0.35281892 0.50311715 0.77145899
```

```
alpha=0.05
```

```
sum(coh.chlaXaccndvi$accndvip.ts1<alpha)/nrow(coh.chlaXaccndvi)
```

```
## [1] 0.06666667
```

```
sum(coh.chlaXmaxndvi$maxndvip.ts1<alpha)/nrow(coh.chlaXmaxndvi)
```

```
## [1] 0.05925926
```

```
print(cbind(coh.chlaXaccndvi$lagoslakeid, coh.chlaXaccndvi$accndvip.ts1)[coh.chlaXaccndvi$accndvip.ts1<
```

```
##          [,1]      [,2]
## [1,]    5104 0.00249975
## [2,]    5288 0.03719628
## [3,]    6199 0.00659934
## [4,]    6399 0.03289671
## [5,]    6973 0.02009799
## [6,]    7810 0.01519848
## [7,]   79457 0.04749525
## [8,]  136680 0.04729527
## [9,]    5453 0.02519748
```

```

print(cbind(coh.chlaXaccndvi$lagoslakeid, coh.chlaXaccndvi$accndvip.ts2)[coh.chlaXaccndvi$accndvip.ts2<=
##          [,1]      [,2]
## [1,]      249 0.02429757
## [2,]     6301 0.02169783
## [3,]    136466 0.00999900
## [4,]     14815 0.00859914
## [5,]    102115 0.04649535
## [6,]      3280 0.03449655
## [7,]      5463 0.03979602

cor(coh.chlaXaccndvi$accndvicoh.ts1,coh.chlaXaccndvi$accndvicoh.ts2)

## [1] -0.002969988

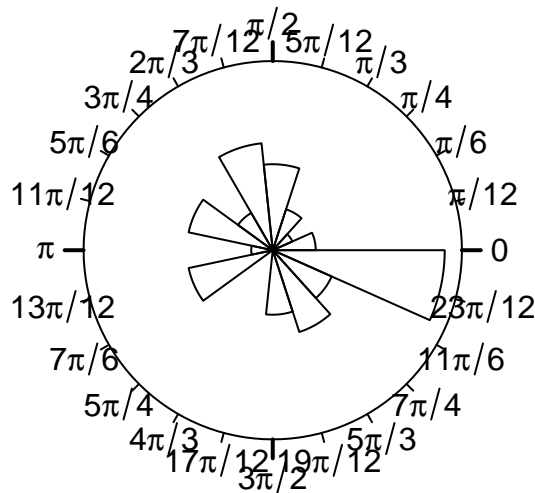
# print(coh.chlaXaccndvi$accndviphi.ts1[coh.chlaXaccndvi$accndvip.ts1<alpha]/pi) #only pattern is that
# print(coh.chlaXmaændvi$maændviphi.ts1[coh.chlaXmaændvi$maændvip.ts1<alpha]/pi)

phicls<-c(-1,-.75,-0.25,0.25,0.75,1)

# hist(coh.chlaXaccndvi$accndviphi.ts1[coh.chlaXaccndvi$accndvip.ts1<0.2]/pi, main="Accumulated NDVI, s
rose(coh.chlaXaccndvi$accndviphi.ts1[coh.chlaXaccndvi$accndvip.ts1<0.3], unit="radian",
     breaks=seq(0,2*pi,length.out=16))

```

**coh.chlaXaccndvi\$accndviphi.ts1[coh.chlaXaccndvi\$accndvip.ts1 <**

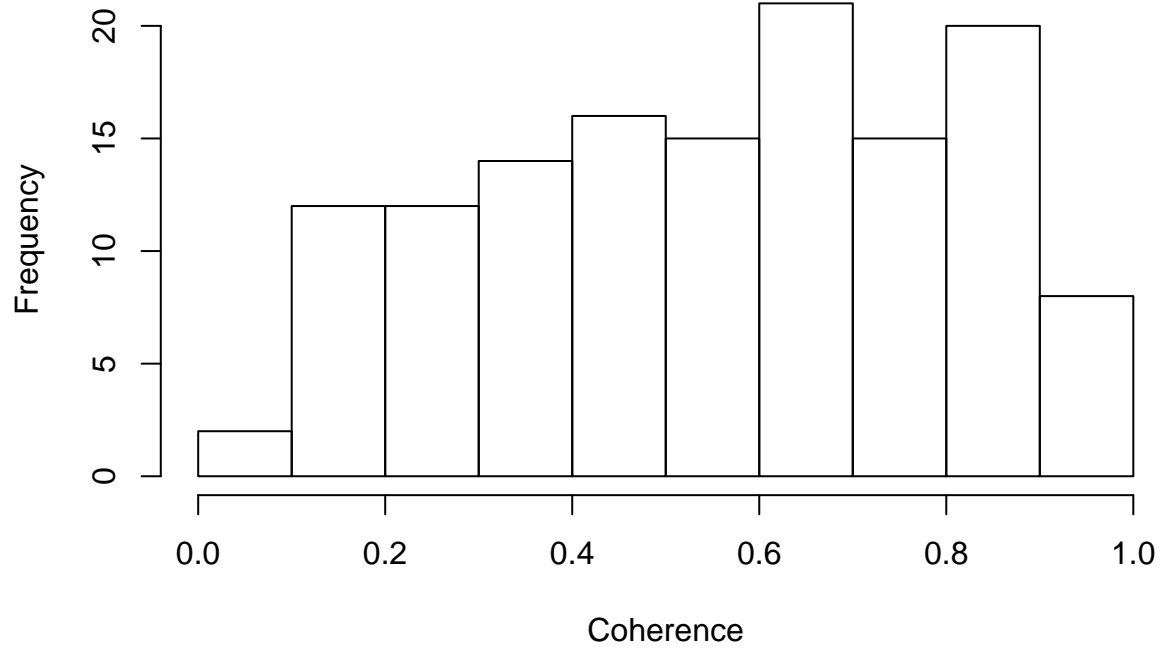


```

#hist(coh.chlaXmaændvi$maændviphi.ts1[coh.chlaXmaændvi$maændvip.ts1<0.2]/pi, main="Maximum NDVI, short
#long timescales
hist(coh.chlaXaccndvi$accndvicoh.ts2, main="Accumulated NDVI, long timescales", xlab="Coherence", ylab=

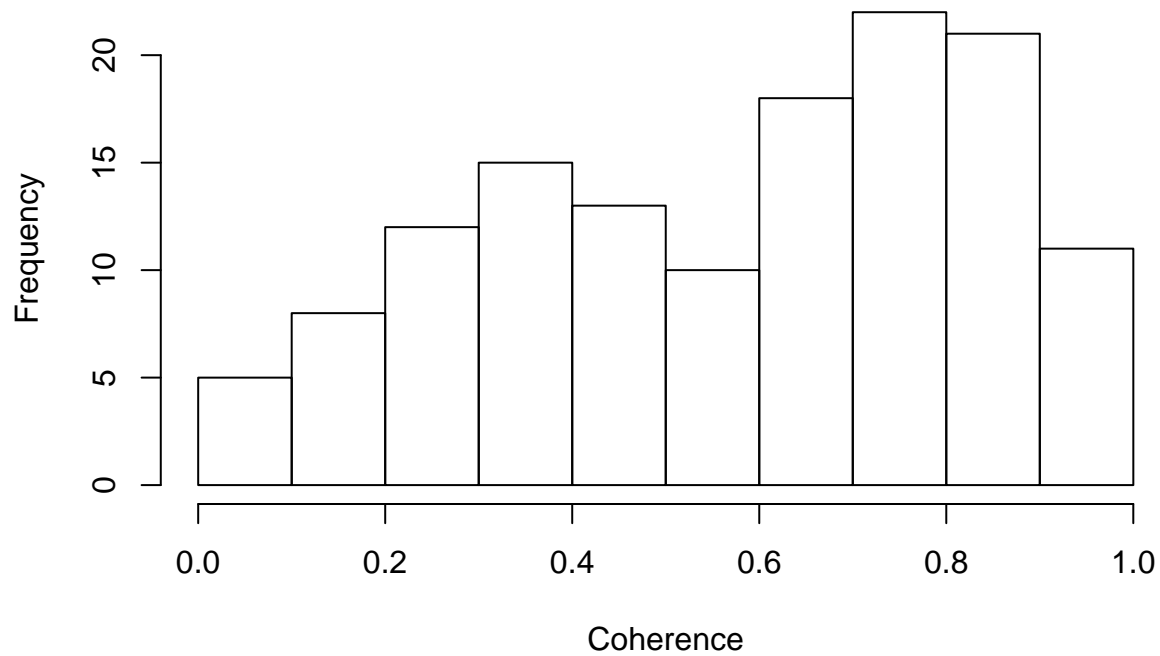
```

### Accumulated NDVI, long timescales



```
hist(coh.chlaXmaxndvi$maxndvicoh.ts2, main="Maximum NDVI, long timescales", xlab="Coherence", ylab="Frequency")
```

### Maximum NDVI, long timescales



```
quantile(coh.chlaXaccndvi$accndvicoh.ts2)
```

```
##          0%          25%          50%          75%         100%
## 0.06700155 0.35635453 0.56072757 0.75753276 0.96052338
```

```
quantile(coh.chlaXmaxndvi$maxndvicoh.ts2)
```

```
##           0%           25%           50%           75%          100%
## 0.04123391 0.35832298 0.61507443 0.78760333 0.96402244
```

```
alpha=0.05
```

```
sum(coh.chlaXaccndvi$accndvip.ts2<alpha)/nrow(coh.chlaXaccndvi)
```

```
## [1] 0.05185185
```

```
sum(coh.chlaXmaxndvi$maxndvip.ts2<alpha)/nrow(coh.chlaXmaxndvi)
```

```
## [1] 0.05185185
```

```
print(coh.chlaXaccndvi$accndviphi.ts2[coh.chlaXaccndvi$accndvip.ts2<alpha]/pi)
```

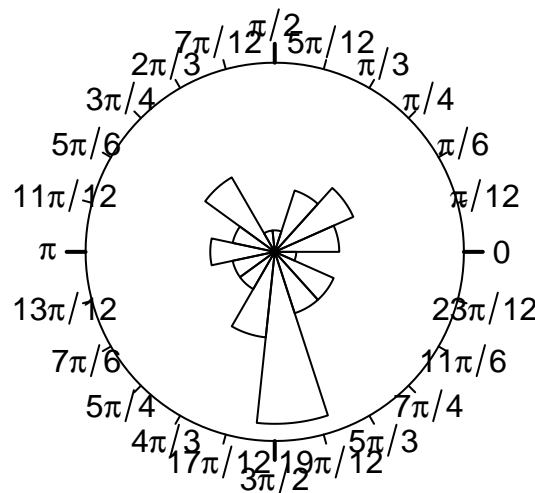
```
## [1] -0.43893809  0.25315167 -0.04386325 -0.65597599 -0.58346970  0.36382168
## [7]  0.89471121
```

```
print(coh.chlaXmaxndvi$maxndviphi.ts2[coh.chlaXmaxndvi$maxndvip.ts2<alpha]/pi)
```

```
## [1] -0.97179292 -0.04190360  0.02097044 -0.67004320 -0.58501674 -0.31373024
## [7] -0.33804686
```

```
# hist(coh.chlaXaccndvi$accndviphi.ts2[coh.chlaXaccndvi$accndvip.ts2<0.2]/pi, main="Accumulated NDVI, l
rose(coh.chlaXaccndvi$accndviphi.ts2[coh.chlaXaccndvi$accndvip.ts2<0.3], unit="radian",
     breaks=seq(0,2*pi,length.out=16))
```

**coh.chlaXaccndvi\$accndviphi.ts2[coh.chlaXaccndvi\$accndvip.ts2 <**



```
#hist(coh.chlaXmaxndvi$maxndviphi.ts1[coh.chlaXmaxndvi$maxndvicoh.ts2>0.6]/pi, main="Maximum NDVI, shor
```

```
states<-readOGR("~/Box Sync/NSF EAGER Synchrony/Data/statesp020.shp")
```

```
## OGR data source with driver: ESRI Shapefile
```

```
## Source: "/Users/jonathanwalter/Box Sync/NSF EAGER Synchrony/Data/statesp020.shp", layer: "statesp020"
```

```
## with 2895 features
```

```
## It has 9 fields
```

```
## Integer64 fields read as strings: STATESP020 DAY_ADM YEAR_ADM
```

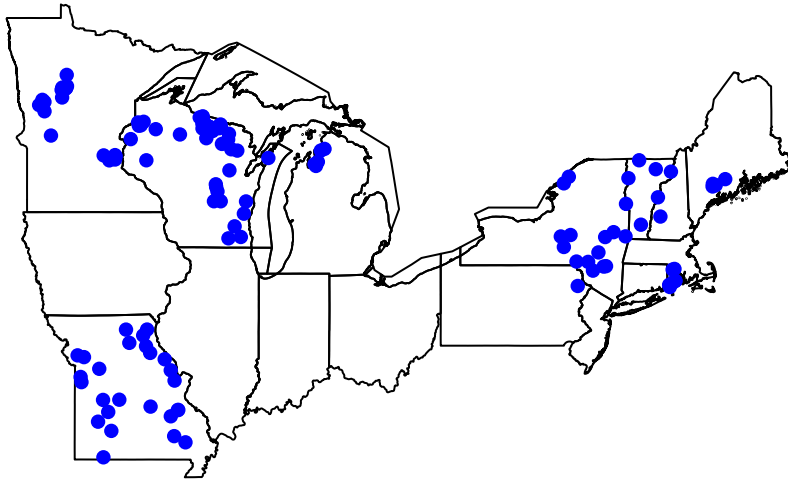
```

getstates<-c("Minnesota", "Iowa", "Wisconsin", "Illinois", "Missouri", "Michigan", "Indiana", "Ohio", "Pennsylvania")
lagosstates<-states[states@data$STATE %in% getstates,]

plot(lagosstates, main="Lakes selected for analysis")
points(analysislakes$lakeinfo$nhd_long, analysislakes$lakeinfo$nhd_lat, pch=16, cex=1, col="blue")

```

## Lakes selected for analysis



```

cohplotdata<-left_join(analysislakes$lakeinfo, coh.chlaXaccndvi, by="lagoslakeid")

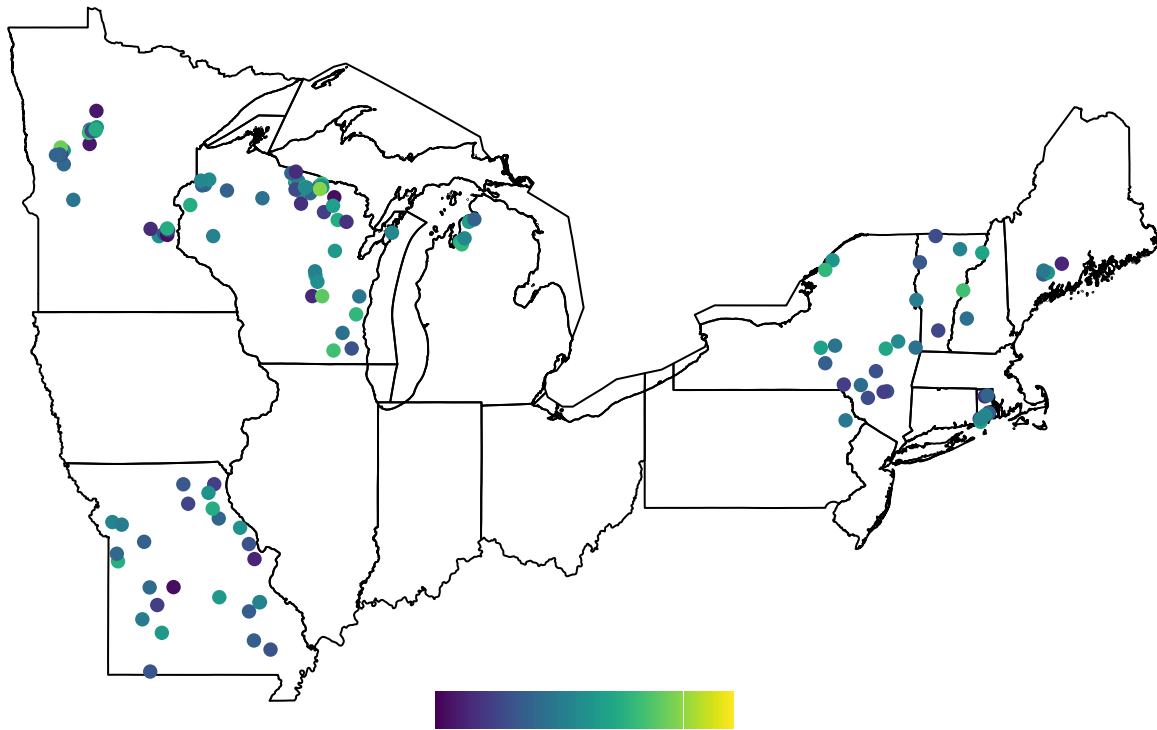
pal<-viridis(100)

par(mar=c(1,0,2,0))

plot(lagosstates, main="Lakes by short timescale coherence")
points(cohplotdata$nhd_long, cohplotdata$nhd_lat, pch=16, cex=1, col=pal[round(cohplotdata$accndvicoh.t
colorbar.plot(x=mean(par("usr")[1:2]),y=par("usr")[3],strip=1:100,col=pal,horizontal = T)

```

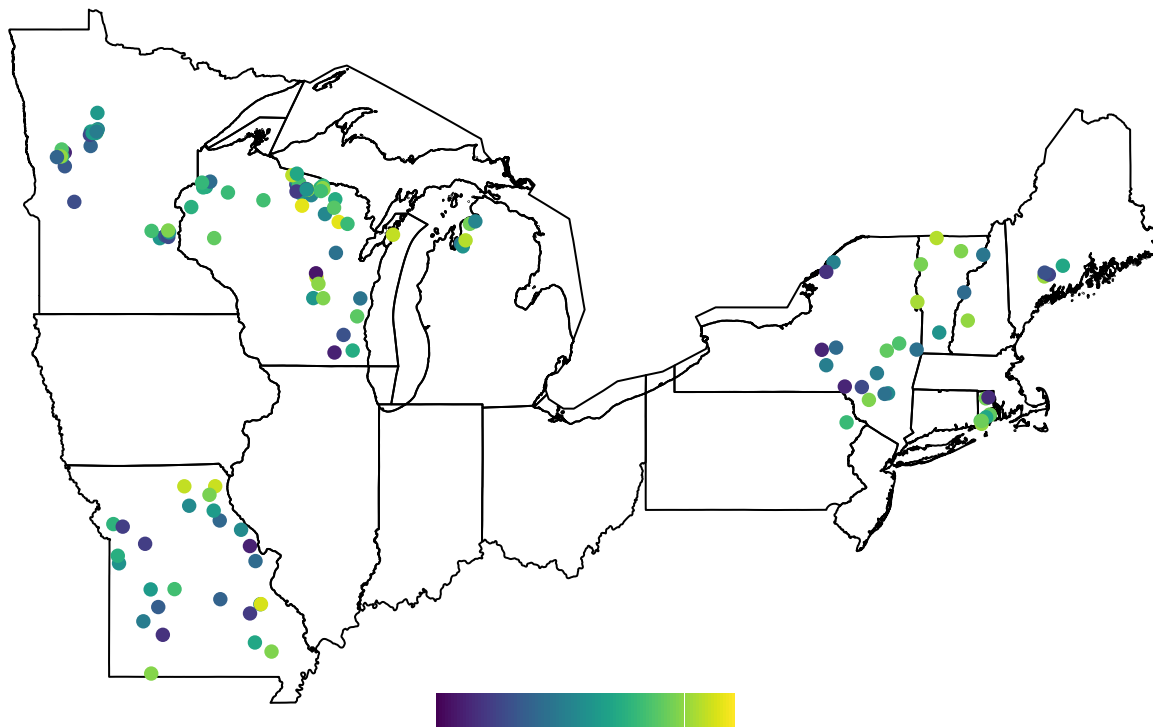
## Lakes by short timescale coherence



```
plot(lagosstates, main="Lakes by long timescale coherence")
points(cohplotdata$nhd_long, cohplotdata$nhd_lat, pch=16, cex=1, col=pal[round(cohplotdata$accndvicoh.t
colorbar.plot(x=mean(par("usr")[1:2]),y=par("usr")[3],strip=1:100,col=pal,horizontal = T)
```



## Lakes by long timescale coherence



*#Need to add: depth, average growing season Chlorophyll-a, TSI(chla) categories, pct ag*

*#agriculture -- is 500m buffer best? Other options include 100m buffer (probably too small) and hu12 wa*  
*# pct.ag<-lagosne\_select(table="buffer500m.lulc", vars=c("lagoslakeid", "buffer500m\_nlcd2001\_pct\_82", "bu*  
 pct.ag<-lagosne\_select(table="hu12.lulc", vars=c("hu12\_zoneid", "hu12\_nlcd2001\_pct\_82", "hu12\_nlcd2006\_pct\_82",  
 pct.ag<-pct.ag[pct.ag\$hu12\_zoneid %in% analysislakes\$lakeinfo\$hu12\_zoneid,]  
 pct.ag.avg<-data.frame(hu12\_zoneid=pct.ag\$hu12\_zoneid, pct.ag=rowMeans(pct.ag[,2:4]))

*#Wetlands*

pct.wetlands<-lagosne\_select(table="hu12.lulc", vars=c(c("hu12\_zoneid", "hu12\_nlcd2001\_pct\_90", "hu12\_nlcd2006\_pct\_90",  
 "hu12\_nlcd2011\_pct\_90", "hu12\_nlcd2001\_pct\_95",  
 "hu12\_nlcd2011\_pct\_95"))))  
 pct.wetlands<-pct.wetlands[pct.wetlands\$hu12\_zoneid %in% analysislakes\$lakeinfo\$hu12\_zoneid,]  
 pct.wetlands\$sum2001<-rowSums(pct.wetlands[,c(2,5)])  
 pct.wetlands\$sum2006<-rowSums(pct.wetlands[,c(3,6)])  
 pct.wetlands\$sum2011<-rowSums(pct.wetlands[,c(4,7)])  
 pct.wetlands.avg<-data.frame(hu12\_zoneid=pct.wetlands\$hu12\_zoneid, pct.wetlands=rowMeans(pct.wetlands[,c(2,5,6,7)]))

*#depth*

depth<-lagosne\_select(table="lakes\_limno", vars=c("lagoslakeid", "maxdepth"))  
 depth<-depth[depth\$lagoslakeid %in% analysislakes\$lakeinfo\$lagoslakeid,] *#use max depth because it's more accurate*

*#growing season Chlorophyll-a*

chla<-lagosne\_select(table="epi\_nutr", vars=c("lagoslakeid", "samplemonth", "chla"))  
 chla<-chla[chla\$lagoslakeid %in% analysislakes\$lakeinfo\$lagoslakeid,]  
 gs.chla<-chla[chla\$samplemonth %in% 5:9,]  
 avg.chla<-aggregate(chla ~ lagoslakeid, data=gs.chla, FUN=mean, na.rm=T)

```

#growing season DOC
doc<-lagosne_select(table="epi_nutr", vars=c("lagoslakeid", "samplemonth", "doc"))
doc<-doc[doc$lagoslakeid %in% analysislakes$lakeinfo$lagoslakeid,]
gs.doc<-doc[doc$samplemonth %in% 5:9,]
avg.doc<-aggregate(doc ~ lagoslakeid, data=gs.doc, FUN=mean, na.rm=T)

#Chlorophyll-a TSI class
#TSI(CHL) = 9.81 ln(CHL) + 30.6
tsi.chl<-data.frame(lagoslakeid=avg.chla$lagoslakeid, tsi=9.81 * log(avg.chla$chla) + 30.6)
tsi.chl$tsi.cat<-rep("lake", nrow(tsi.chl))

tsi.chl$tsi.cat[tsi.chl$tsi < 40]<-"oligotrophic"
tsi.chl$tsi.cat[tsi.chl$tsi >=40 & tsi.chl$tsi < 50]<-"mesotrophic"
tsi.chl$tsi.cat[tsi.chl$tsi >=50 & tsi.chl$tsi < 70]<-"eutrophic"
tsi.chl$tsi.cat[tsi.chl$tsi >= 70] <-"hypereutrophic"

#CV of terrestrial NDVI
cv.accndvi<-NULL
for(lake in 1:length(analysislakes$lakedata)){
  tmp<-analysislakes$lakedata[[lake]][rownames(analysislakes$lakedata[[lake]])=="accndvi",]
  cv.accndvi<-c(cv.accndvi, sd(tmp)/mean(tmp))
  # rm(tmp)
}
cv.accndvi<-data.frame(lagoslakeid=as.numeric(names(analysislakes$lakedata)), cv.accndvi=cv.accndvi)

#mean precipitation
prcp.normal<-raster("~/Box Sync/NSF EAGER Synchrony/Data/PRISM Data/PRISM_ppt_30yr_normal_800mM2_annual_
lakepts<-SpatialPoints(coords=cbind(analysislakes$lakeinfo$nhd_long, analysislakes$lakeinfo$nhd_lat))
lake.prcp<-data.frame(lagoslakeid=analysislakes$lakeinfo$lagoslakeid, prcp.normal=raster::extract(prcp.

#huc2 and huc4 watershed codes
huc_codes<-read.csv("/Users/jonathanwalter/GitHub/AquaTerrSynch/AnalysisCode/match_huc_codes.csv", colC

#state info
states<-lagosne_select(table="state", vars=c("state_zoneid", "state_name"))

#shoreline development ratio
sdev<-analysislakes$lakeinfo$lake_perim_meters/(2*sqrt(pi*analysislakes$lakeinfo$lake_area_ha*10000))
shoredev<-data.frame(lagoslakeid=analysislakes$lakeinfo$lagoslakeid, shoredev=sdev)

predictors<-analysislakes$lakeinfo
predictors<-left_join(predictors, depth, by="lagoslakeid")
predictors<-left_join(predictors, pct.ag.avg, by="hu12_zoneid")

## Warning: Column `hu12_zoneid` joining factors with different levels,
## coercing to character vector
predictors<-left_join(predictors, pct.wetlands.avg, by="hu12_zoneid")

## Warning: Column `hu12_zoneid` joining character vector and factor, coercing
## into character vector
predictors<-left_join(predictors, avg.chla, by="lagoslakeid")
predictors<-left_join(predictors, tsi.chl, by="lagoslakeid")
predictors<-left_join(predictors, states, by="state_zoneid")

```

```
## Warning: Column `state_zoneid` joining factors with different levels,
## coercing to character vector
```

```
predictors<-left_join(predictors, cv.acndvi, by="lagoslakeid")
predictors<-left_join(predictors, avg.doc, by="lagoslakeid")
predictors<-left_join(predictors, lake.prcp, by="lagoslakeid")
predictors<-left_join(predictors, shoredev, by="lagoslakeid")
#predictors<-left_join(predictors, huc_codes, by="hu4_zoneid")

for(nn in 1:ncol(predictors)){

  if(is.factor(predictors[,nn])){
    predictors[,nn]<-factor(predictors[,nn])
  }

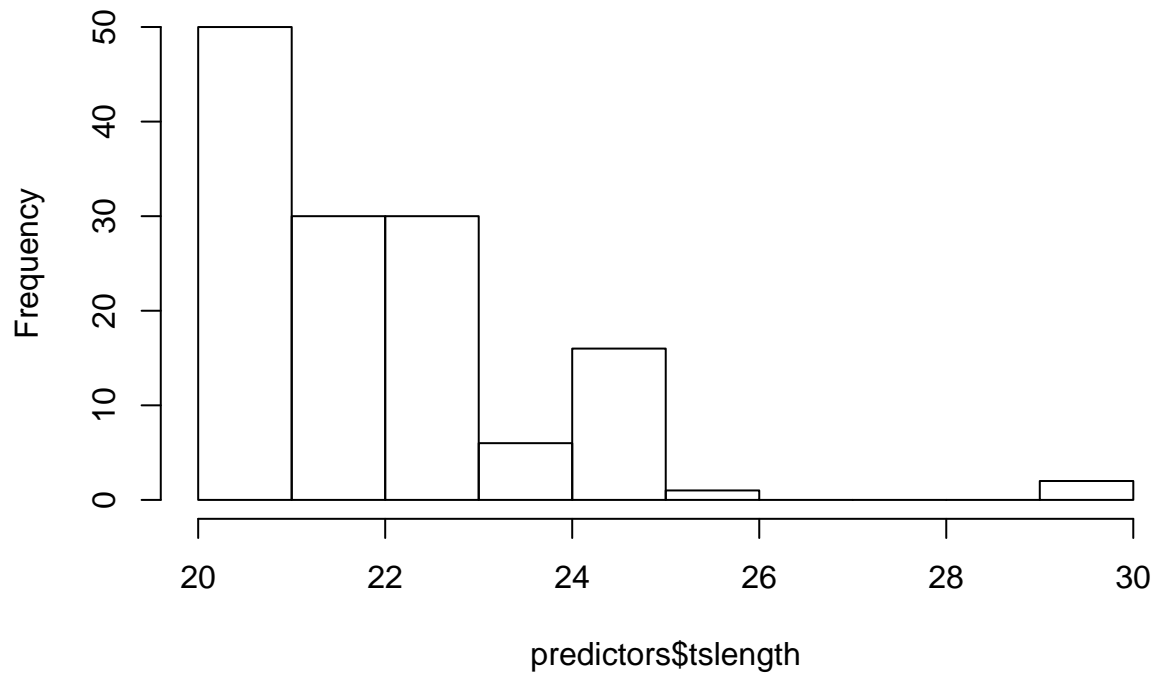
}

str(predictors)
```

```
## 'data.frame': 135 obs. of 26 variables:
## $ lagoslakeid : num 211 249 618 906 969 ...
## $ gnis_name : chr NA NA "Butternut Lake" "Sparkling Lake" ...
## $ nhd_lat : num 44.5 43.7 45.9 46 45.8 ...
## $ nhd_long : num -73.3 -73.4 -89 -89.7 -89.3 ...
## $ lake_area_ha : num 113496.4 30 504.7 63.7 210.2 ...
## $ lake_perim_meters: num 1042251 3494 13134 3777 9402 ...
## $ nhd_ftype : int 390 390 390 390 390 390 390 390 390 390 ...
## $ nhd_fcode : int 39004 39004 39004 39004 39004 39004 39004 39004 39004 39004 ...
## $ hu4_zoneid : Factor w/ 28 levels "HU4_10","HU4_12",...: 17 17 11 8 12 10 10 10 10 10 ...
## $ hu12_zoneid : chr "HU12_17646" "HU12_16835" "HU12_13309" "HU12_13098" ...
## $ state_zoneid : chr "State_17" "State_5" "State_9" "State_9" ...
## $ elevation_m : num 28.8 28.2 514.5 494.7 503.3 ...
## $ start : num 1989 1990 1993 1989 1994 ...
## $ end : num 2010 2010 2013 2011 2013 ...
## $ tslength : num 22 21 21 23 20 21 21 21 21 22 ...
## $ maxdepth : num 97 NA 12.8 20 11.6 ...
## $ pct.ag : num 2.5298 0.4199 0.0976 0.3029 6.6886 ...
## $ pct.wetlands : num 5.3 7.27 32.8 19.36 48.32 ...
## $ chla : num 5.39 7.94 2.44 1.86 2.04 ...
## $ tsi : num 47.1 50.9 39.4 36.7 37.6 ...
## $ tsi.cat : chr "mesotrophic" "eutrophic" "oligotrophic" "oligotrophic" ...
## $ state_name : Factor w/ 10 levels "Maine","Michigan",...: 9 6 10 10 10 2 2 2 2 2 ...
## $ cv.acndvi : num 0.0572 0.0542 0.0443 0.0561 0.0417 ...
## $ doc : num 5.07 4.41 NA 3.36 1.46 ...
## $ prcp.normal : num 895 931 794 796 793 ...
## $ shoredev : num 8.73 1.8 1.65 1.34 1.83 ...
```

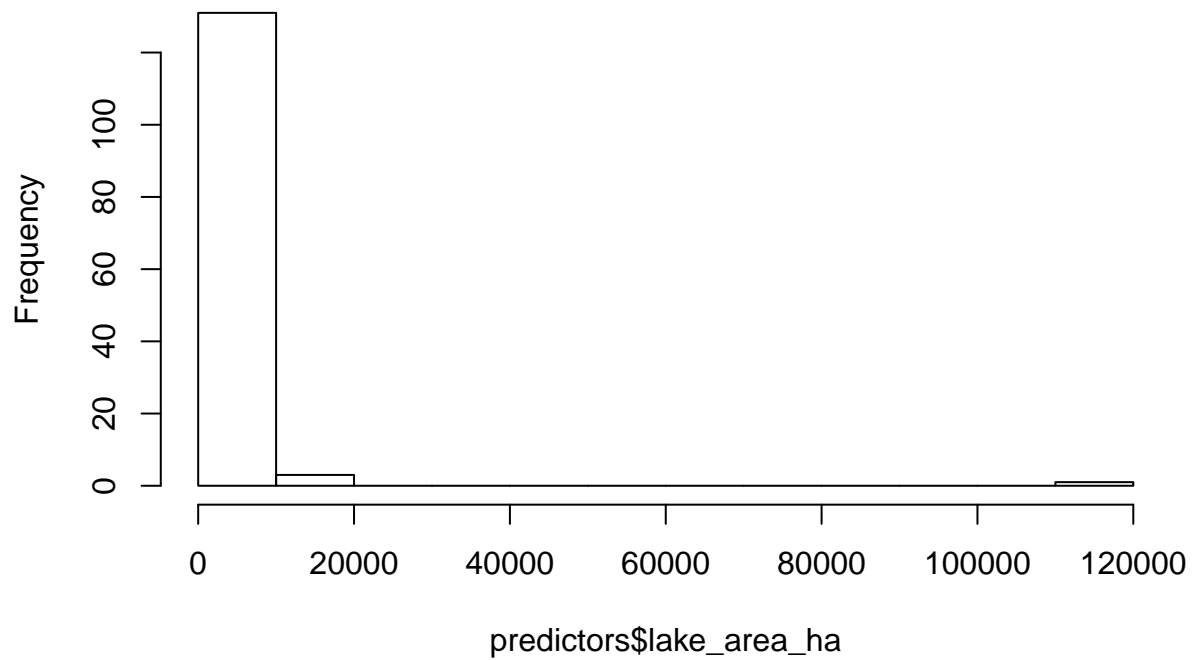
```
hist(predictors$tslength)
```

**Histogram of predictors\$tslength**



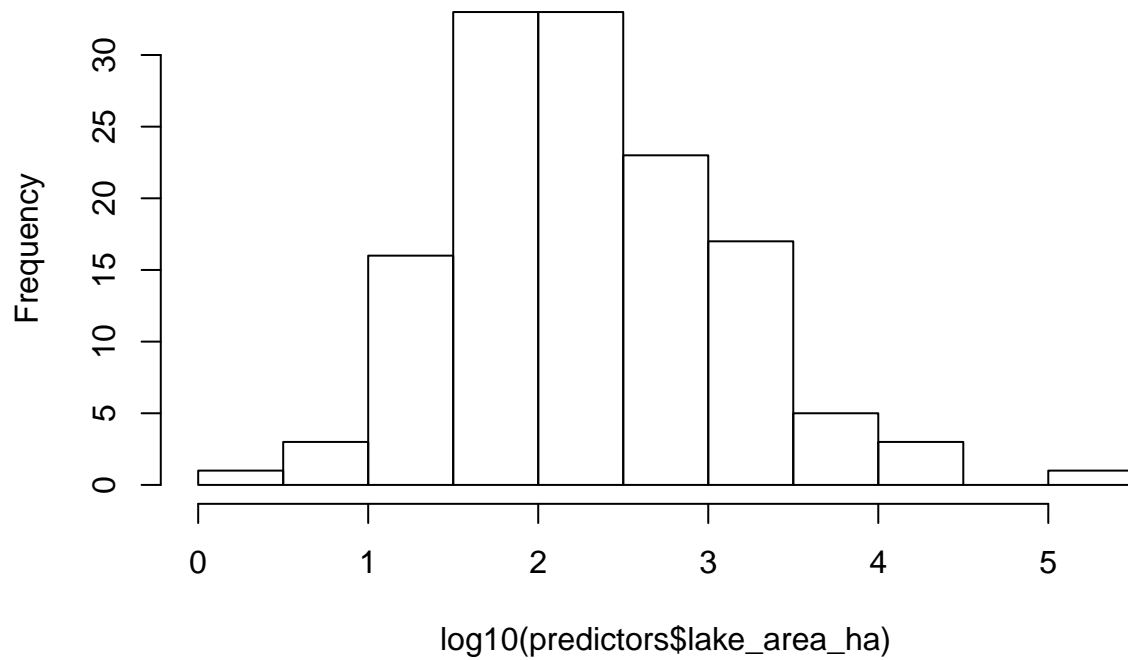
```
hist(predictors$lake_area_ha)
```

**Histogram of predictors\$lake\_area\_ha**



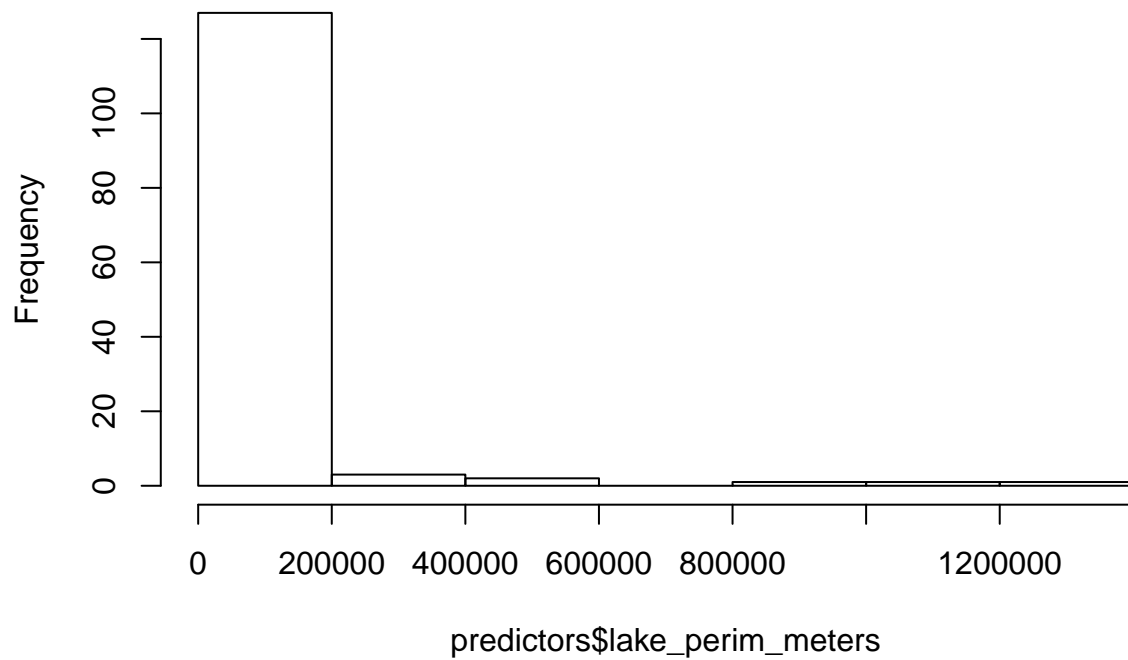
```
hist(log10(predictors$lake_area_ha))
```

**Histogram of  $\log_{10}(\text{predictors\$lake\_area\_ha})$**



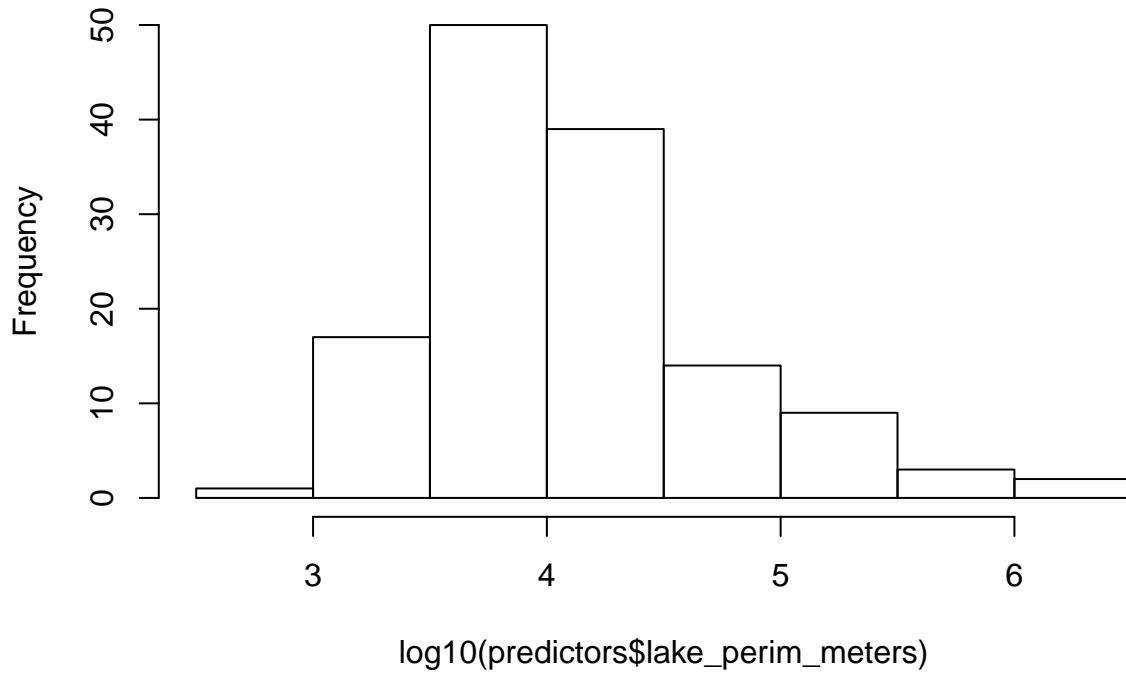
```
hist(predictors$lake_perim_meters)
```

**Histogram of predictors\$lake\_perim\_meters**



```
hist(log10(predictors$lake_perim_meters))
```

## Histogram of log10(predictors\$lake\_perim\_meters)



```
table(predictors$nhd_fcode)
```

```
##
## 39000 39004 39009 39010 39012 43601
##      1   110    14     3     6     1
```

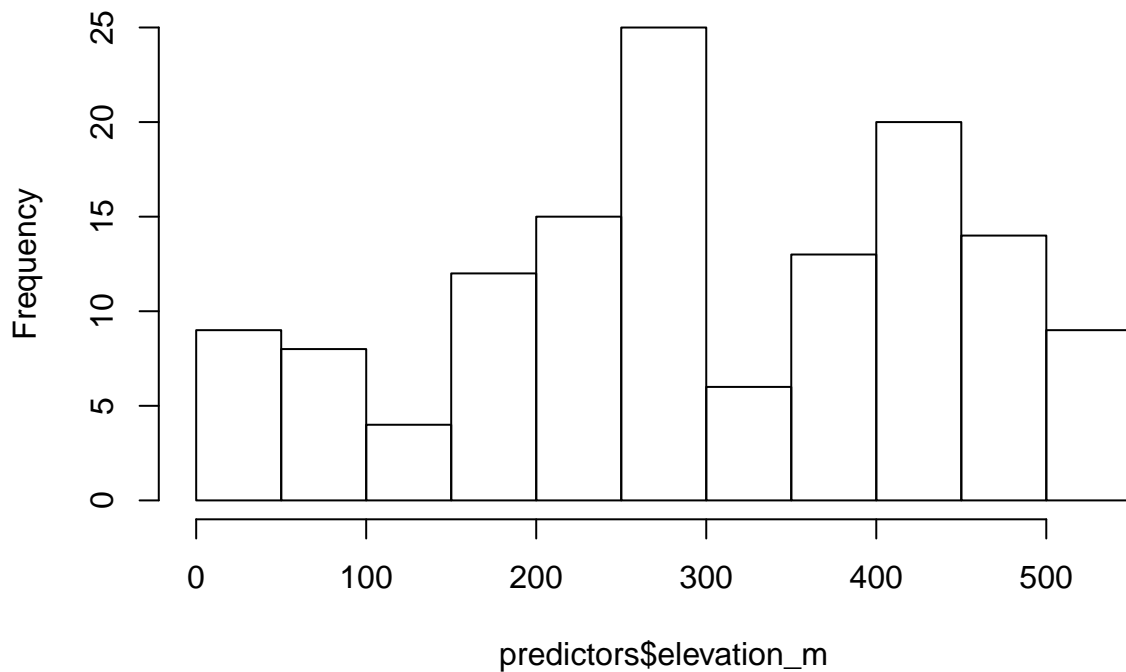
```
table(predictors$hu12_zoneid)
```

```
##
## HU12_10463 HU12_10471 HU12_10488 HU12_10493 HU12_10499 HU12_10676
##           1           2           1           1           1           1
## HU12_10700 HU12_10785 HU12_10862 HU12_10863 HU12_10865 HU12_11197
##           1           1           1           4           1           1
## HU12_11495 HU12_11509 HU12_11514 HU12_11515 HU12_11522 HU12_11768
##           1           1           2           1           1           1
## HU12_11816 HU12_11826 HU12_11829 HU12_11889 HU12_11938 HU12_11978
##           1           1           1           1           1           1
## HU12_12113 HU12_12125 HU12_12225 HU12_13098 HU12_13100 HU12_13125
##           1           1           1           5           1           1
## HU12_13164 HU12_13192 HU12_13234 HU12_13241 HU12_13244 HU12_13261
##           1           1           1           1           1           1
## HU12_13300 HU12_13304 HU12_13309 HU12_13354 HU12_13360 HU12_13370
##           1           1           2           1           1           3
## HU12_13374 HU12_13376 HU12_13388 HU12_13413 HU12_13616 HU12_13624
##           1           1           1           1           1           1
## HU12_13628 HU12_13633 HU12_13634 HU12_14494 HU12_14495 HU12_14496
##           1           1           1           1           1           1
## HU12_14497 HU12_14533 HU12_148 HU12_1494 HU12_15183 HU12_15280
##           4           1           1           1           1           1
## HU12_15296 HU12_15315 HU12_15329 HU12_1537 HU12_15856 HU12_16122
```

```
##      1      1      1      1      1      1
## HU12_16125 HU12_1615 HU12_1621 HU12_16347 HU12_16746 HU12_16747
##      1      1      2      1      1      2
## HU12_16749 HU12_16835 HU12_16882 HU12_17143 HU12_17178 HU12_17235
##      1      1      1      1      1      2
## HU12_17401 HU12_17407 HU12_17433 HU12_17477 HU12_17488 HU12_17504
##      1      1      1      1      2      2
## HU12_17512 HU12_17513 HU12_17541 HU12_17646 HU12_17651 HU12_17655
##      1      2      1      1      1      1
## HU12_1802 HU12_18174 HU12_1819 HU12_1828 HU12_18730 HU12_1896
##      1      1      1      1      1      1
## HU12_19726 HU12_1980 HU12_19842 HU12_20279 HU12_2173 HU12_2200
##      1      1      1      1      1      1
## HU12_2239 HU12_2410 HU12_2412 HU12_2429 HU12_4337 HU12_4347
##      1      1      1      1      1      1
## HU12_442 HU12_488 HU12_509 HU12_542 HU12_581 HU12_829
##      1      1      1      1      1      1
```

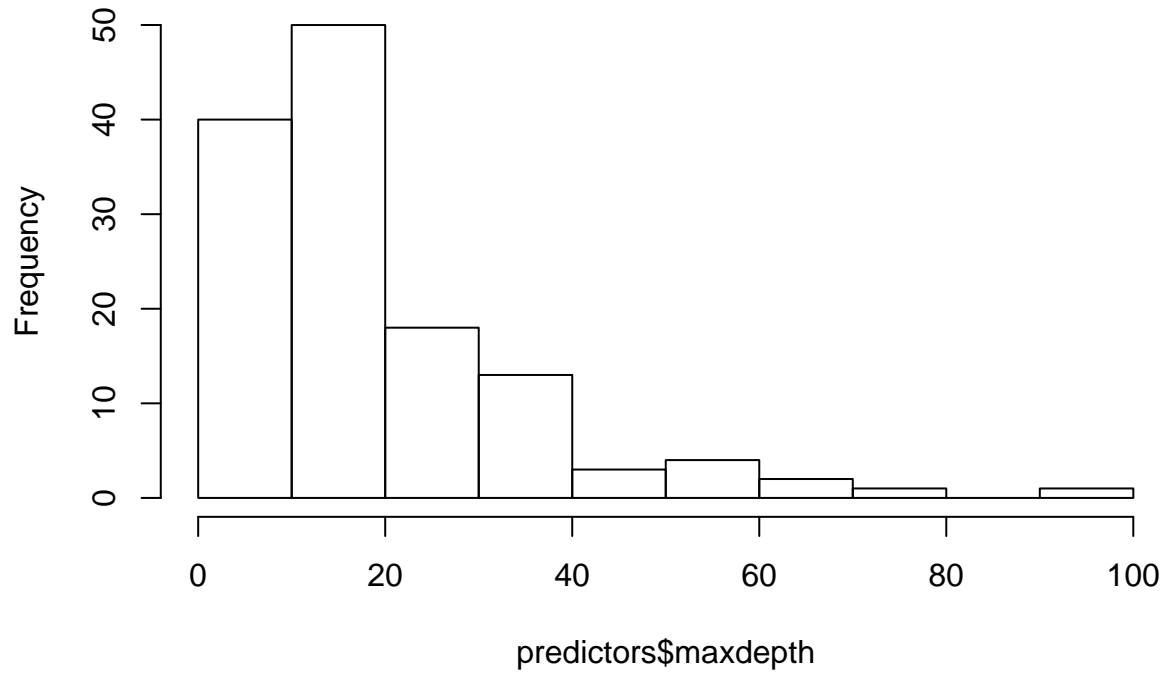
```
hist(predictors$elevation_m)
```

**Histogram of predictors\$elevation\_m**



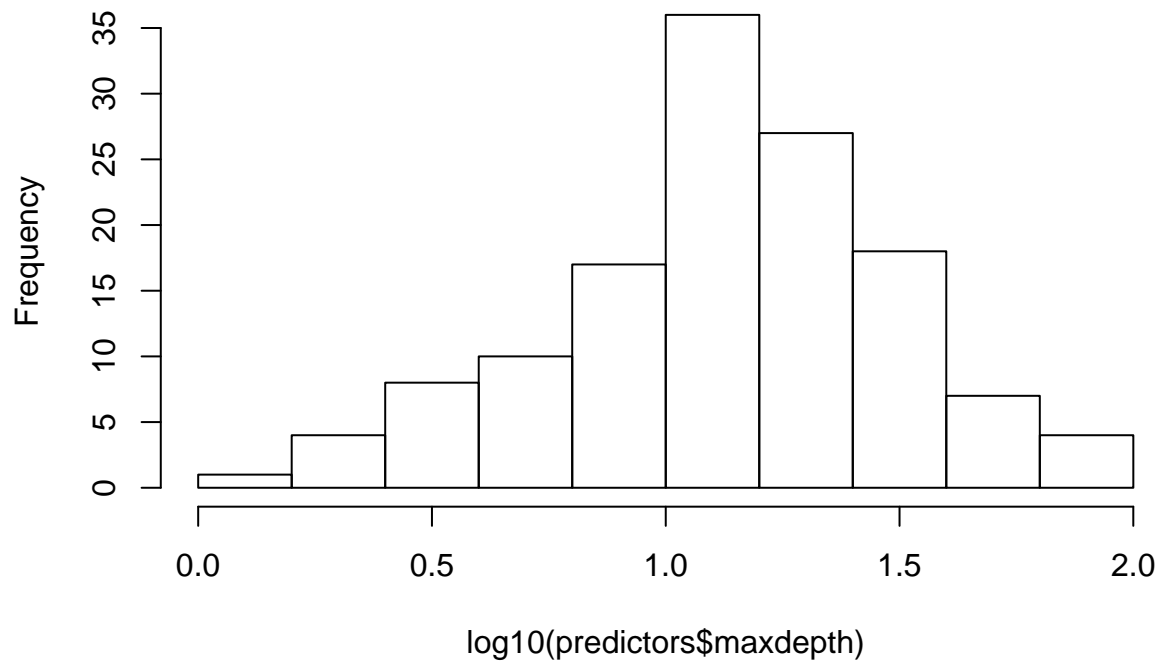
```
hist(predictors$maxdepth)
```

**Histogram of predictors\$maxdepth**



```
hist(log10(predictors$maxdepth))
```

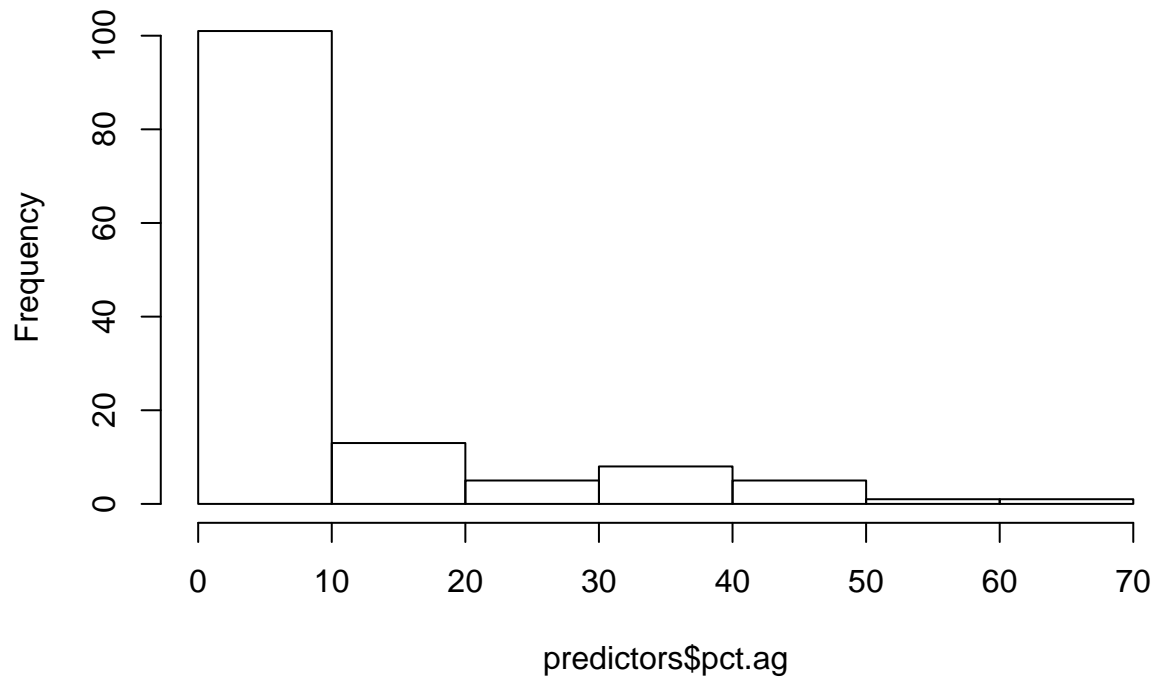
**Histogram of log10(predictors\$maxdepth)**



```
hist(predictors$pct.ag)
```

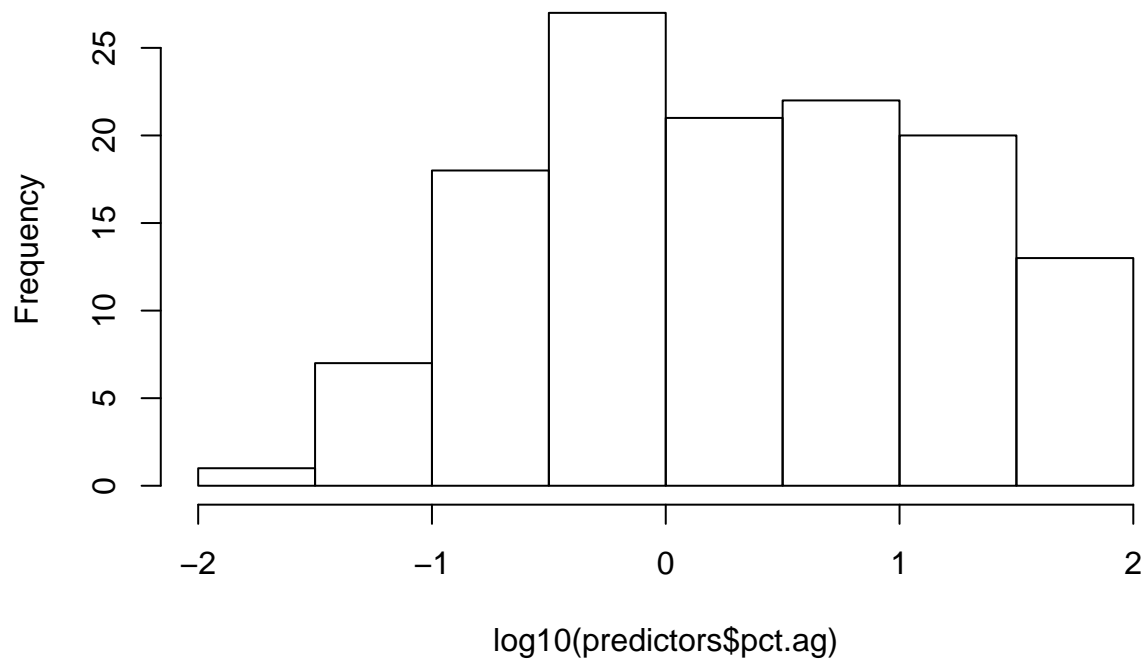


**Histogram of predictors\$pct.ag**



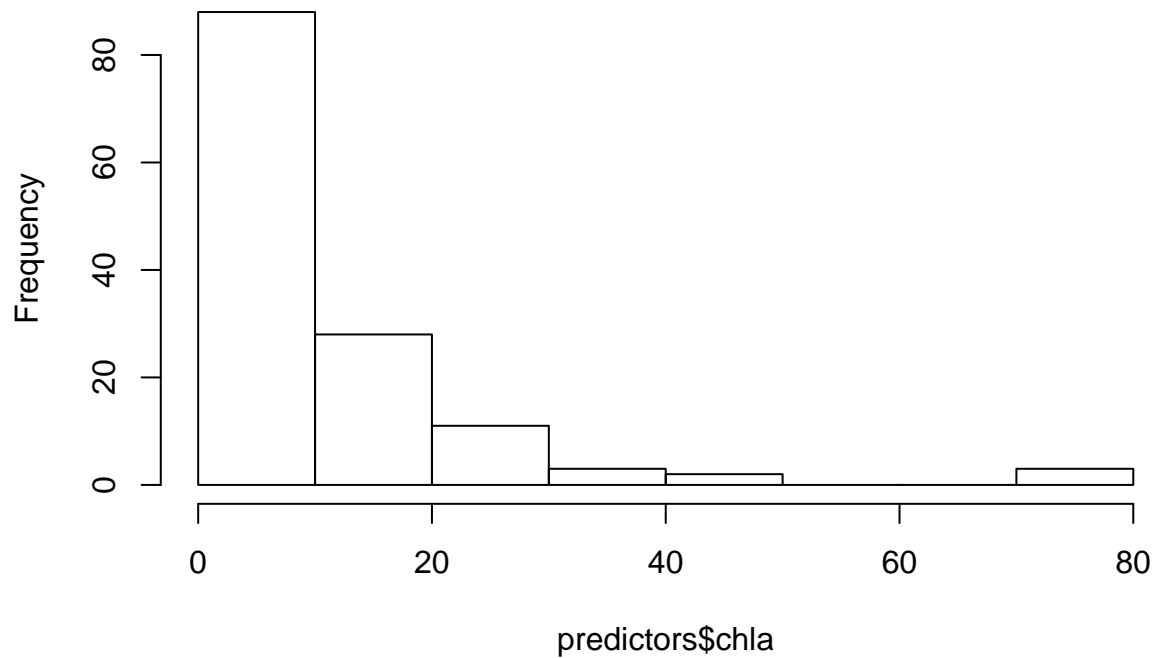
```
hist(log10(predictors$pct.ag))
```

**Histogram of log10(predictors\$pct.ag)**



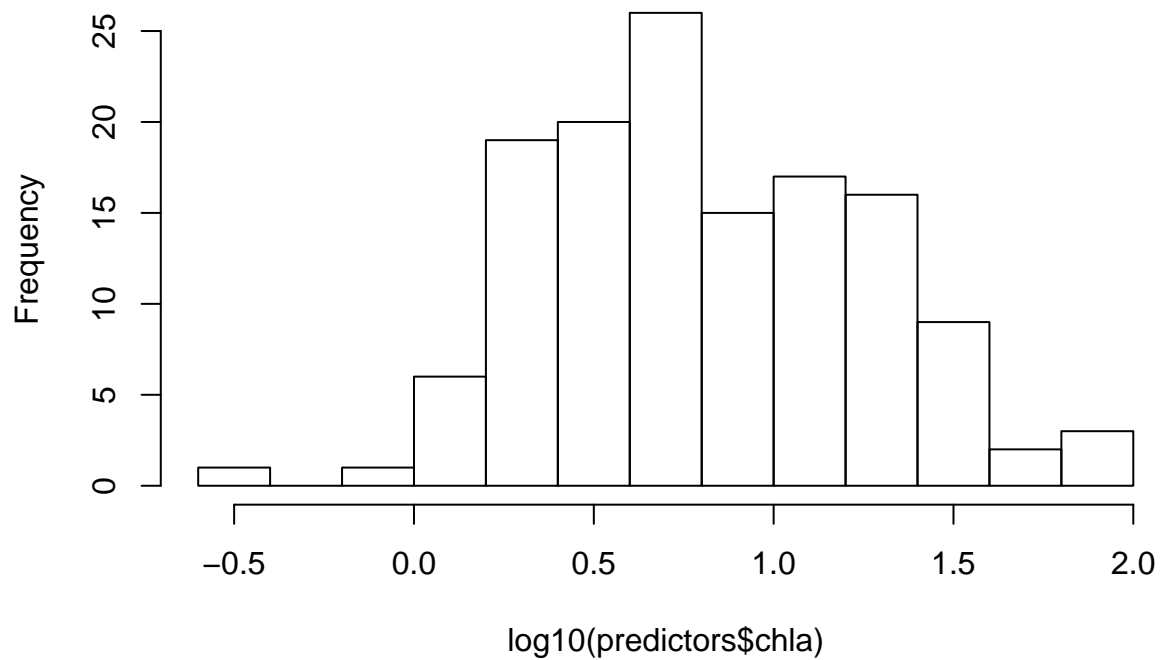
```
hist(predictors$chla)
```

**Histogram of predictors\$chla**



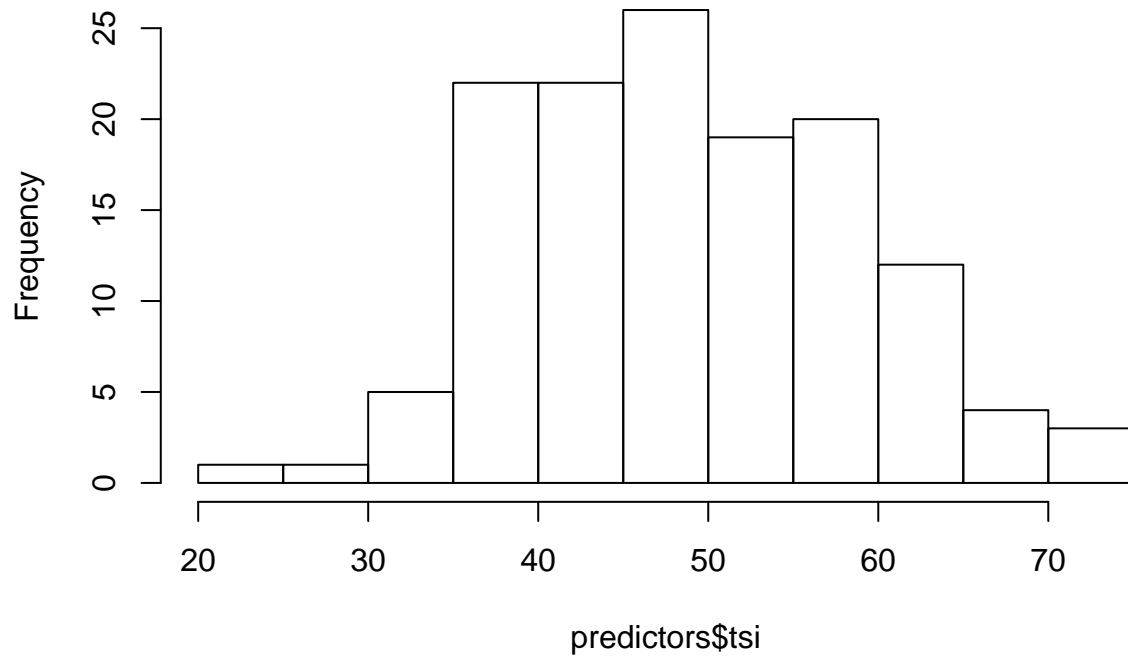
```
hist(log10(predictors$chla))
```

**Histogram of log10(predictors\$chla)**



```
hist(predictors$tsi)
```

## Histogram of predictors\$tsi

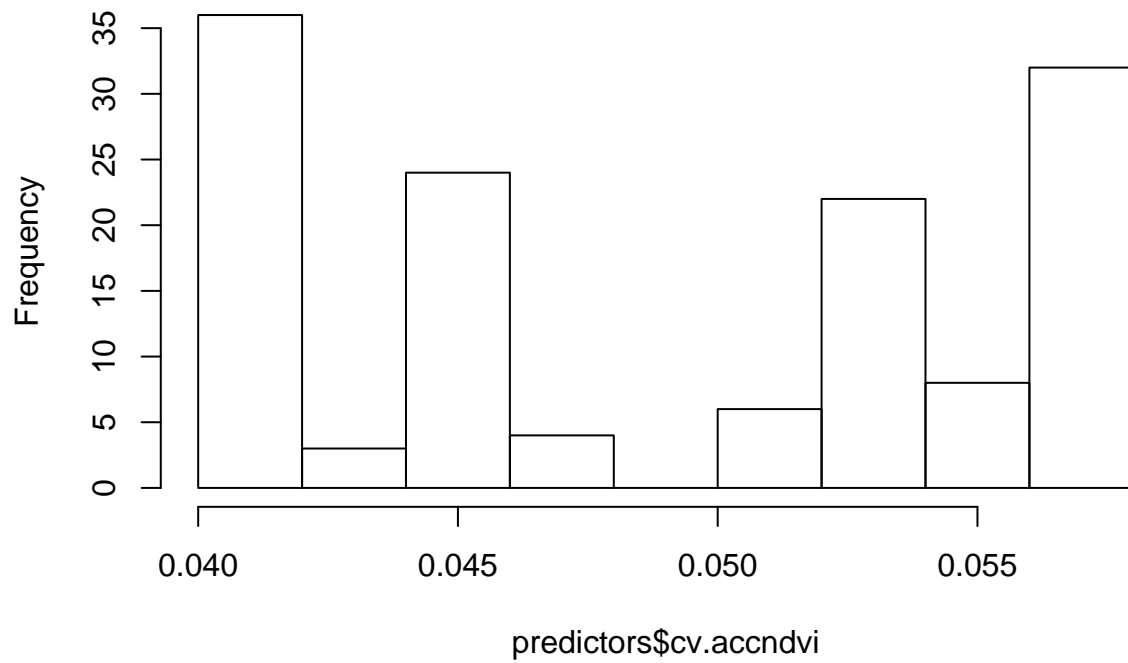


```
table(predictors$tsi.cat)
```

```
##  
##      eutrophic hypereutrophic   mesotrophic  oligotrophic  
##          55           3         48         29
```

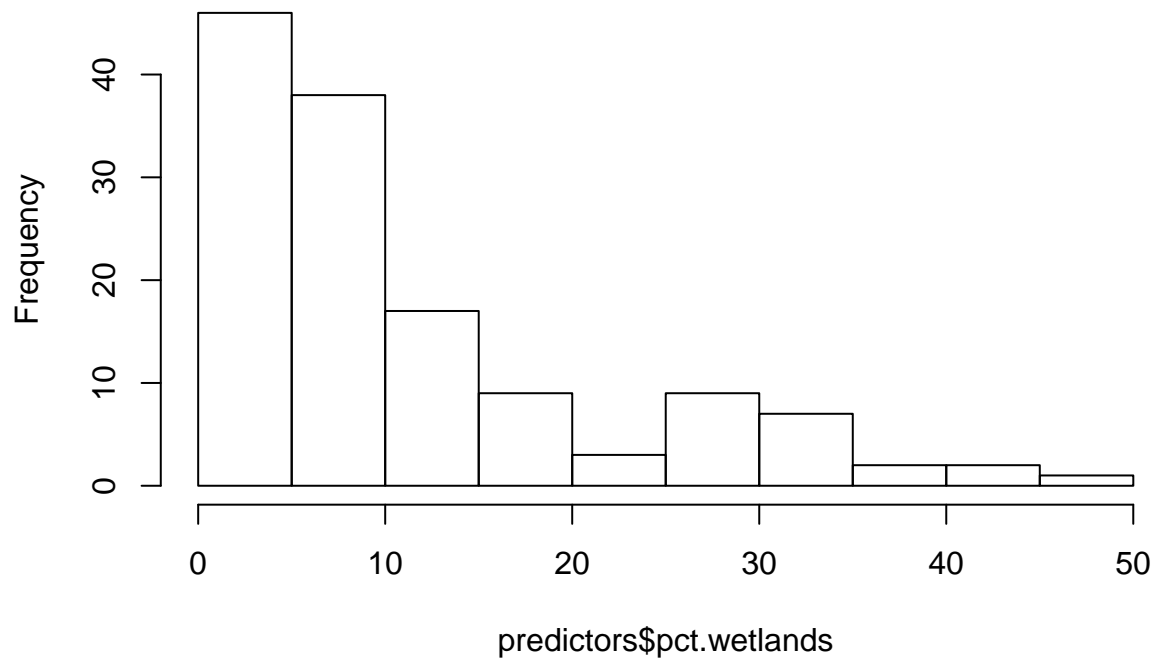
```
hist(predictors$cv.acndvi)
```

**Histogram of predictors\$cv.accndvi**



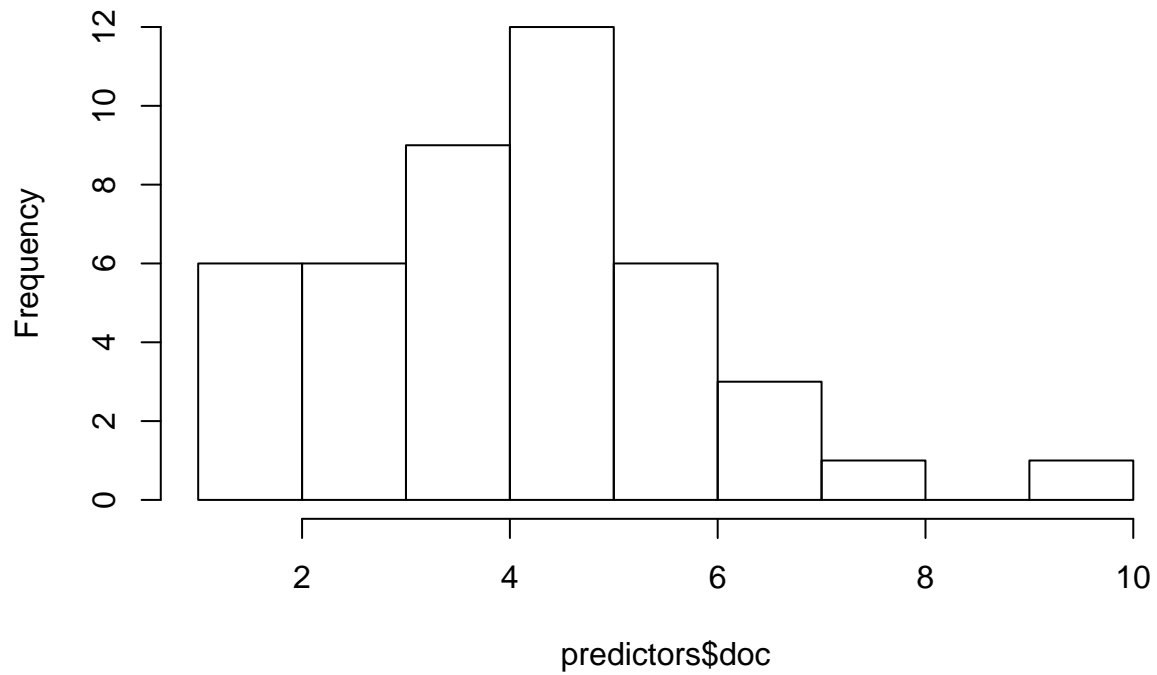
```
hist(predictors$pct.wetlands)
```

**Histogram of predictors\$pct.wetlands**



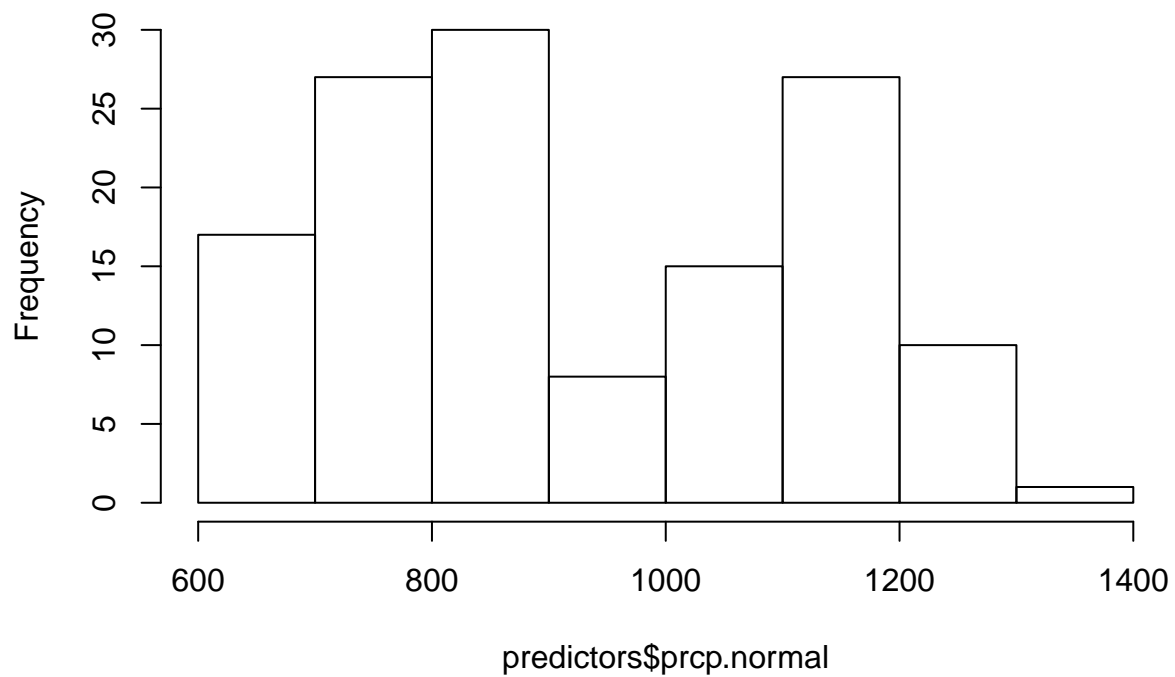
```
hist(predictors$doc)
```

**Histogram of predictors\$doc**



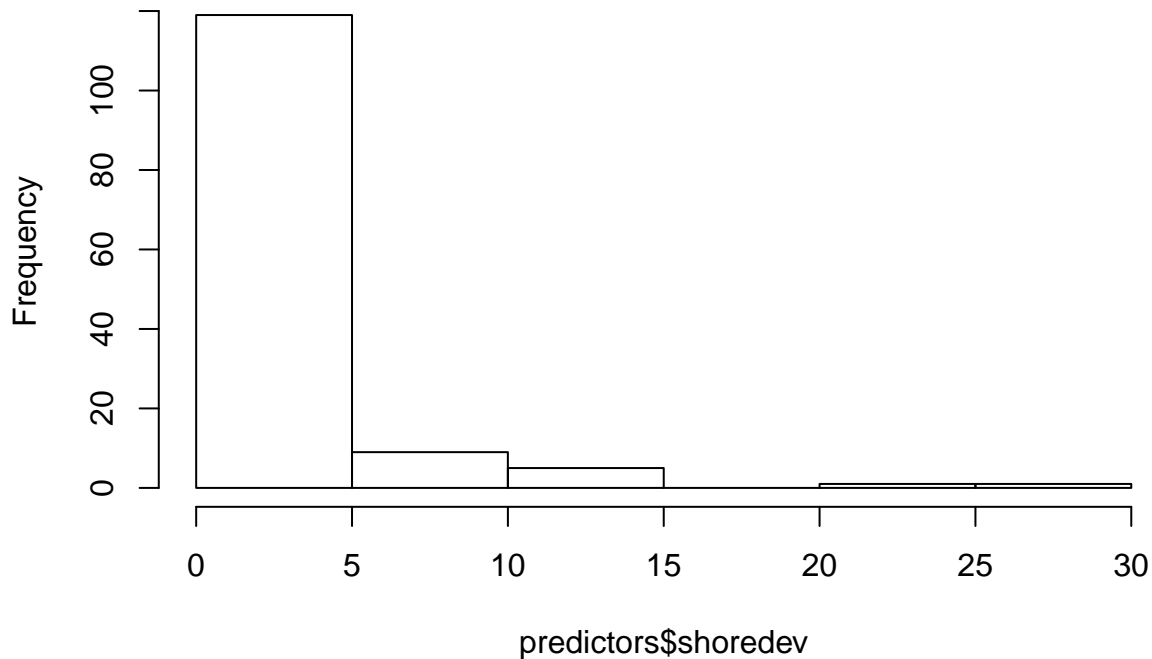
```
hist(predictors$prcp.normal)
```

**Histogram of predictors\$prcp.normal**



```
hist(predictors$shoreddev)
```

## Histogram of predictors\$shoredev



```
# predictors$log10_lake_area_ha<-log10(predictors$lake_area_ha) #not necessary to transform with random
# predictors$log10_lake_perim_meters<-log10(predictors$lake_perim_meters)
# predictors$log10_maxdepth<-log10(predictors$maxdepth)
# predictors$log10_pct.ag<-log10(predictors$pct.ag+1)
# predictors$log10_chla<-log10(predictors$chla)

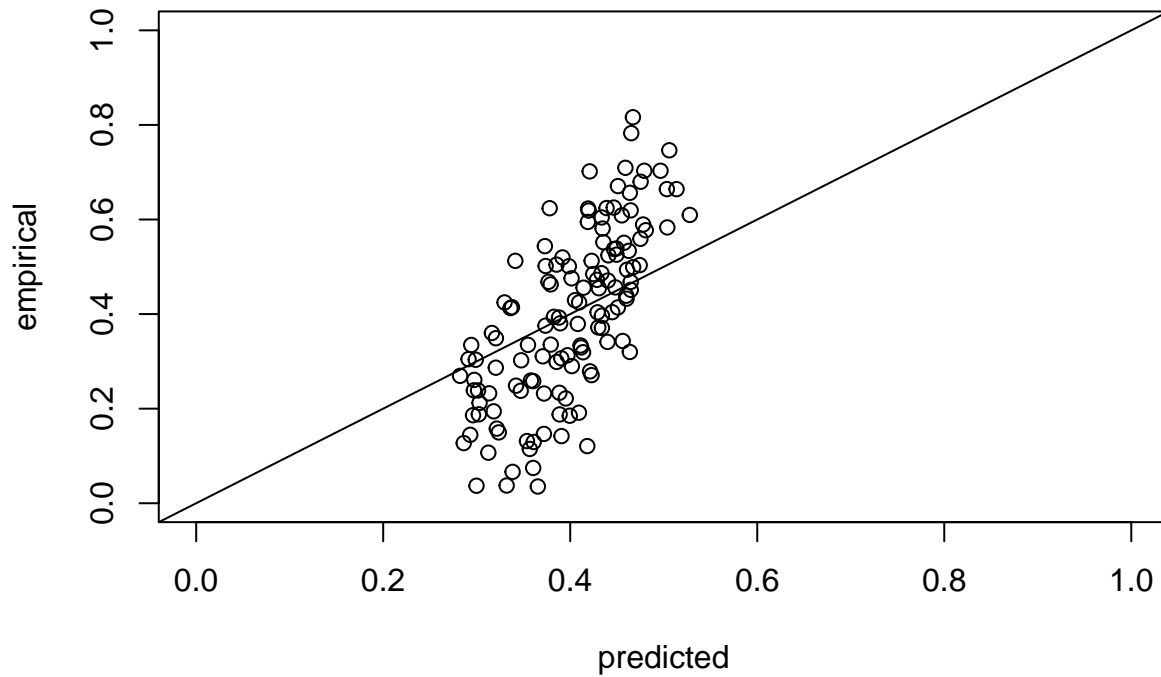
modvars.accndvi<-left_join(predictors, coh.chlaXaccndvi, by="lagoslakeid")
modvars.accndvi$nhd_ftype<-factor(modvars.accndvi$nhd_ftype)
modvars.accndvi$tsi.cat<-factor(modvars.accndvi$tsi.cat)
modvars.accndvi$tslength<-modvars.accndvi$end-modvars.accndvi$start + 1

modvars.accndvi<-modvars.accndvi[!is.na(modvars.accndvi$maxdepth),]
modvars.accndvi<-modvars.accndvi[!is.na(modvars.accndvi$pct.ag),]

modvars.accndvi.phist<-modvars.accndvi[modvars.accndvi$accndvip.ts1<0.3,]
modvars.accndvi.philt<-modvars.accndvi[modvars.accndvi$accndvip.ts2<0.3,]

# cforest.st<-partykit::cforest(accndvicoh.ts1 ~ lake_area_ha + lake_perim_meters + maxdepth + pct.ag +
#                               data=modvars.accndvi, ntree=20000)
cforest.st<-party::cforest(accndvicoh.ts1 ~ shoredev + lake_area_ha + maxdepth + pct.ag + chla + tsi.c
                           hu4_zoneid + cv.accndvi + pct.wetlands + doc + prcp.normal,
                           data=modvars.accndvi, controls=cforest_control(ntree=80000))
predcoh.st<-predict(cforest.st, newdata=modvars.accndvi,type="response")
#hist(predcoh.st)
#hist(modvars.accndvi$accndvicoh.ts1)
plot(predcoh.st, modvars.accndvi$accndvicoh.ts1, xlab="predicted", ylab="empirical", main="Coherence, sl
     xlim=c(0,1), ylim=c(0,1))
abline(a=0,b=1)
```

## Coherence, short ts



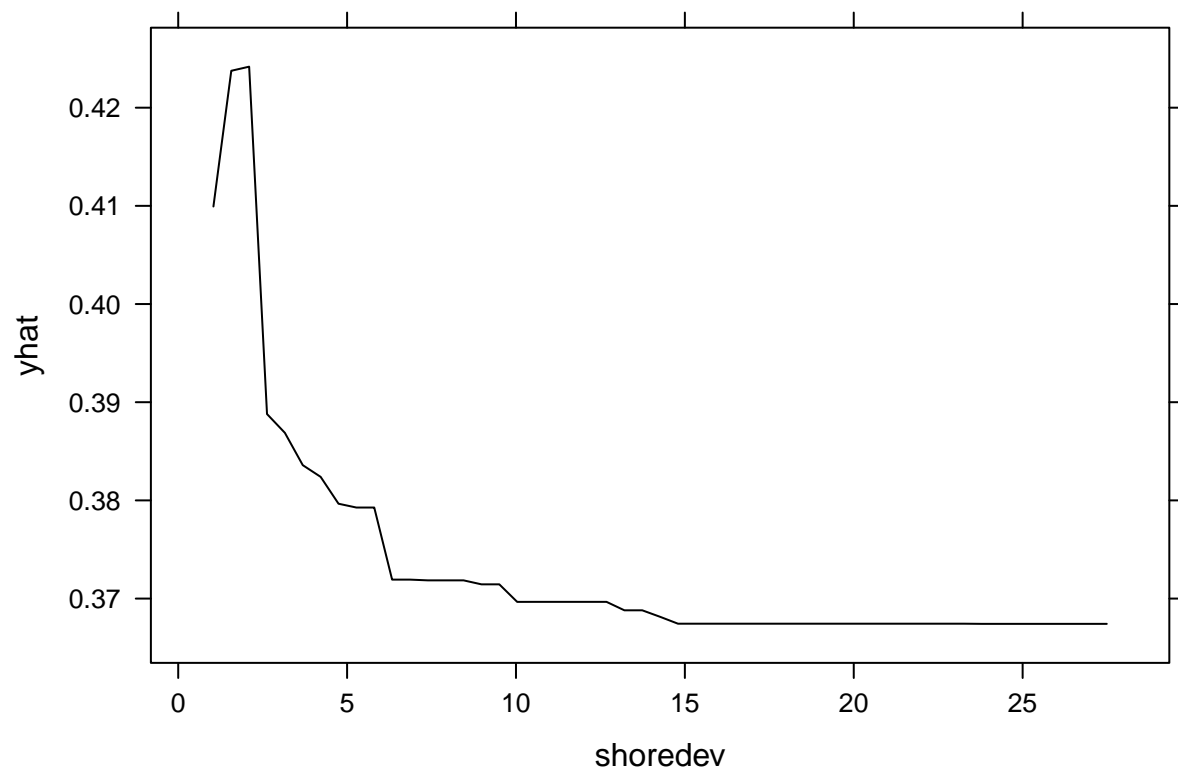
```
cor.test(predcoh.st,modvars.accndvi$accndvicoh.ts1)
```

```
##
## Pearson's product-moment correlation
##
## data: predcoh.st and modvars.accndvi$accndvicoh.ts1
## t = 11.639, df = 129, p-value < 2.2e-16
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.6203220 0.7902151
## sample estimates:
##      cor
## 0.7156966
```

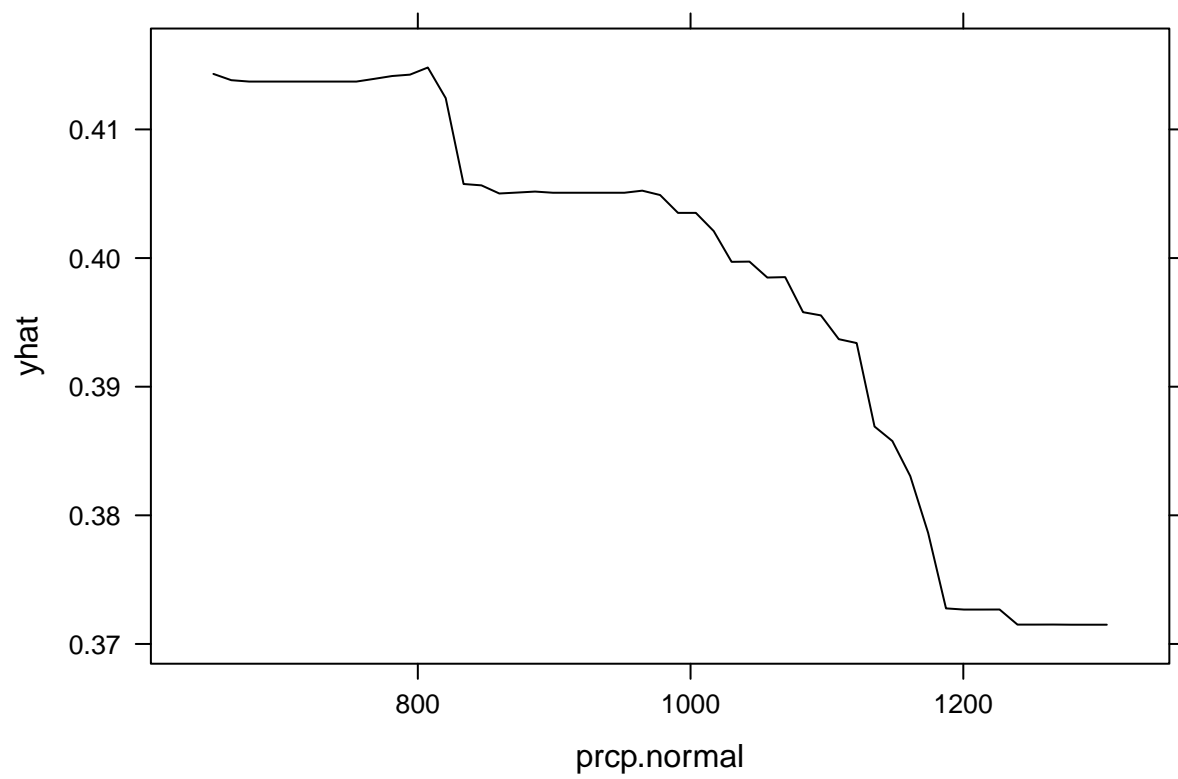
```
varimp.coh.st<-varimp(cforest.st)
print(varimp.coh.st[order(varimp.coh.st, decreasing=T)])
```

```
##      shoredev  prcp.normal      pct.ag      maxdepth  lake_area_ha
## 2.006857e-03  8.283009e-04  4.229068e-04  1.997432e-04  1.771907e-04
##          chla          doc    cv.accndvi  pct.wetlands  hu4_zoneid
## -9.555775e-06 -3.402358e-05 -1.034784e-04 -1.419238e-04 -3.006311e-04
##          tsi.cat
## -4.747599e-04
```

```
partial(cforest.st, pred.var="shoredev", train=modvars.accndvi, type="regression", plot=T)
```

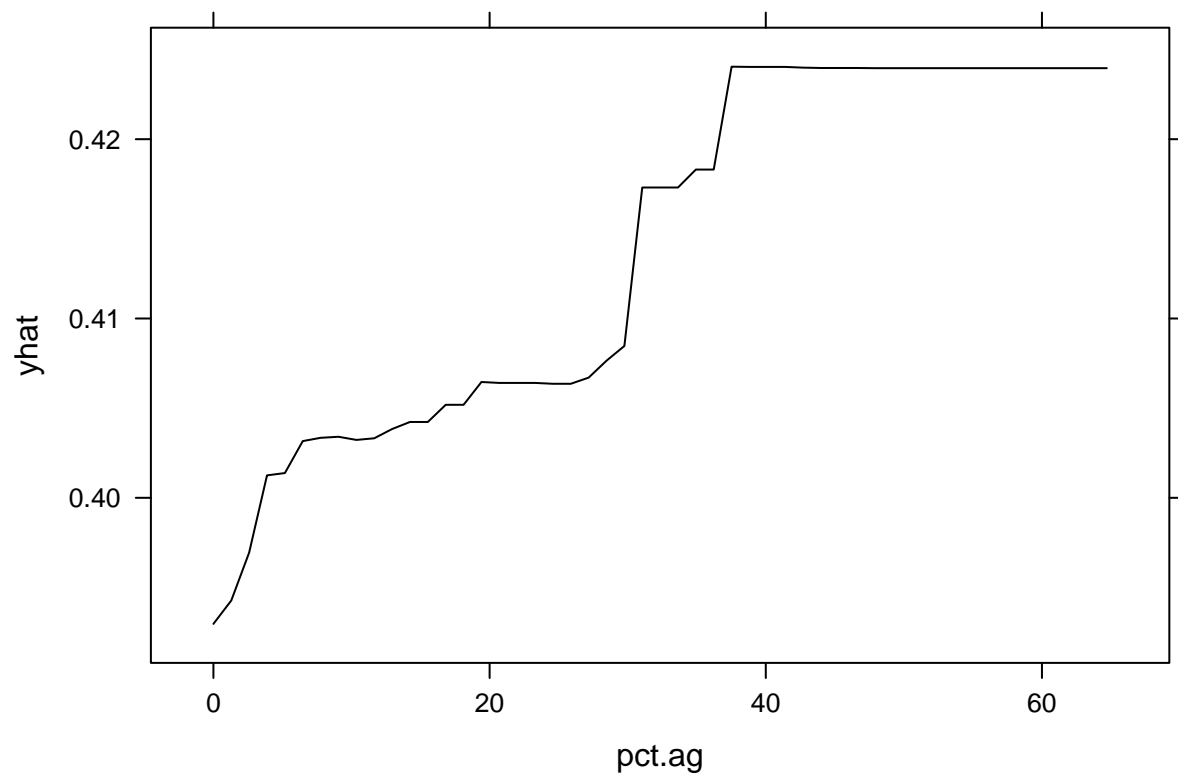


```
partial(cforest.st, pred.var="prcp.normal", train=modvars.accndvi, type="regression", plot=T)
```

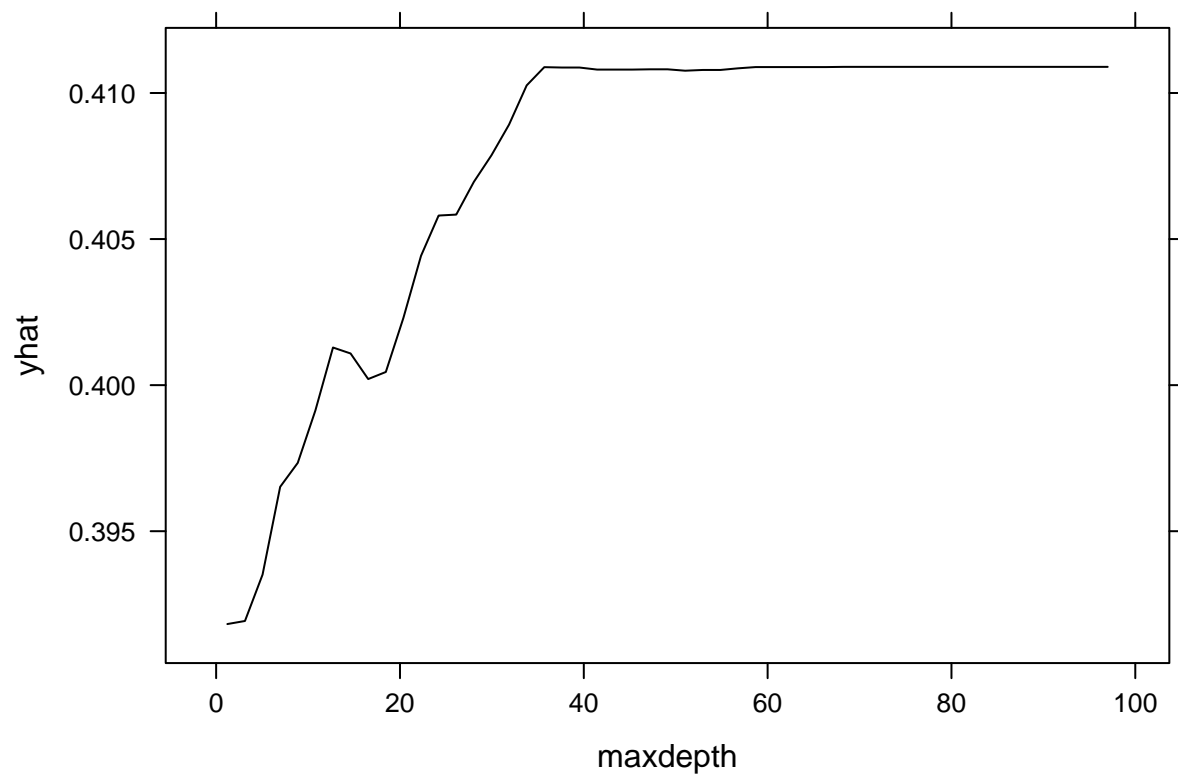


```
partial(cforest.st, pred.var="pct.ag", train=modvars.accndvi, type="regression", plot=T)
```

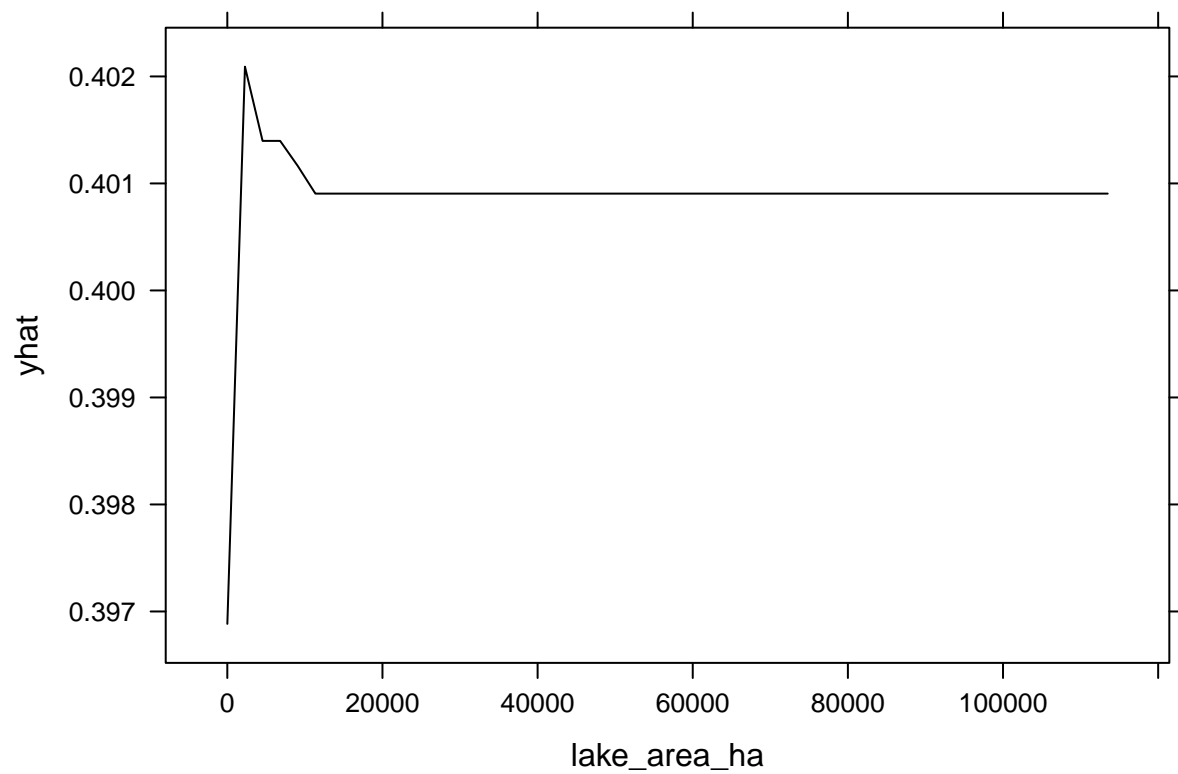




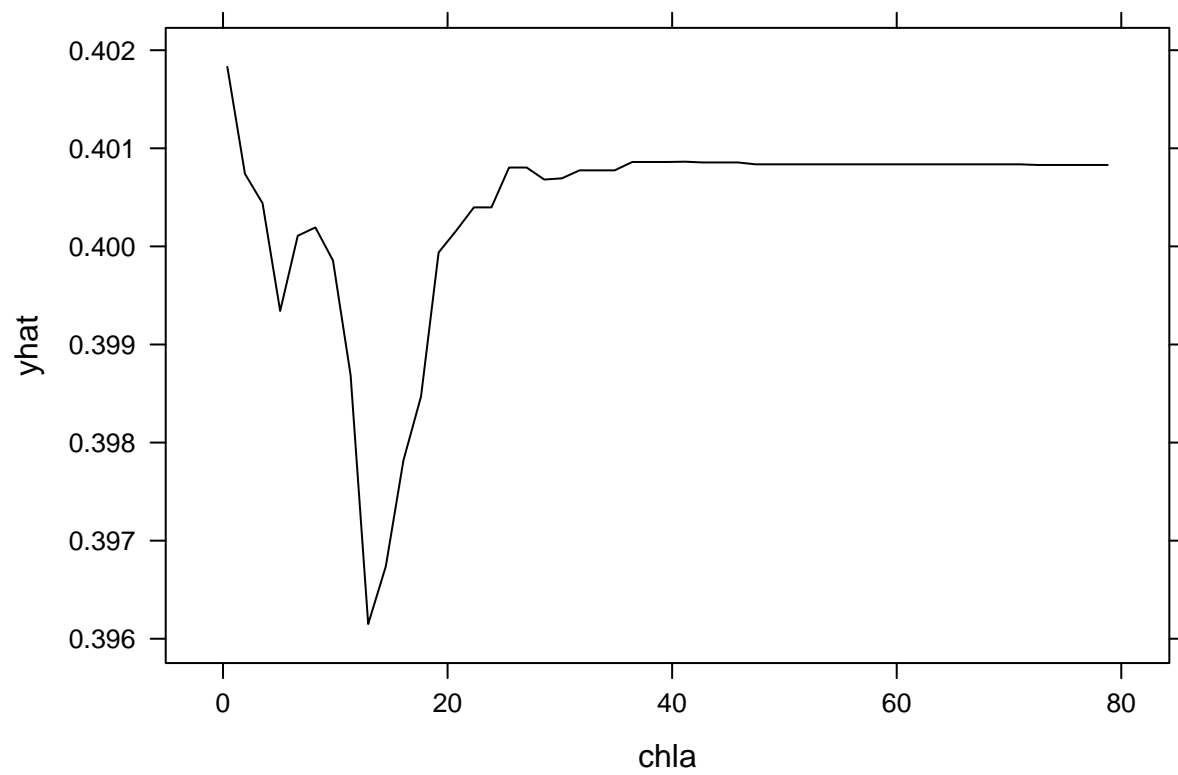
```
partial(cforest.st, pred.var="maxdepth", train=modvars.accndvi, type="regression", plot=T)
```



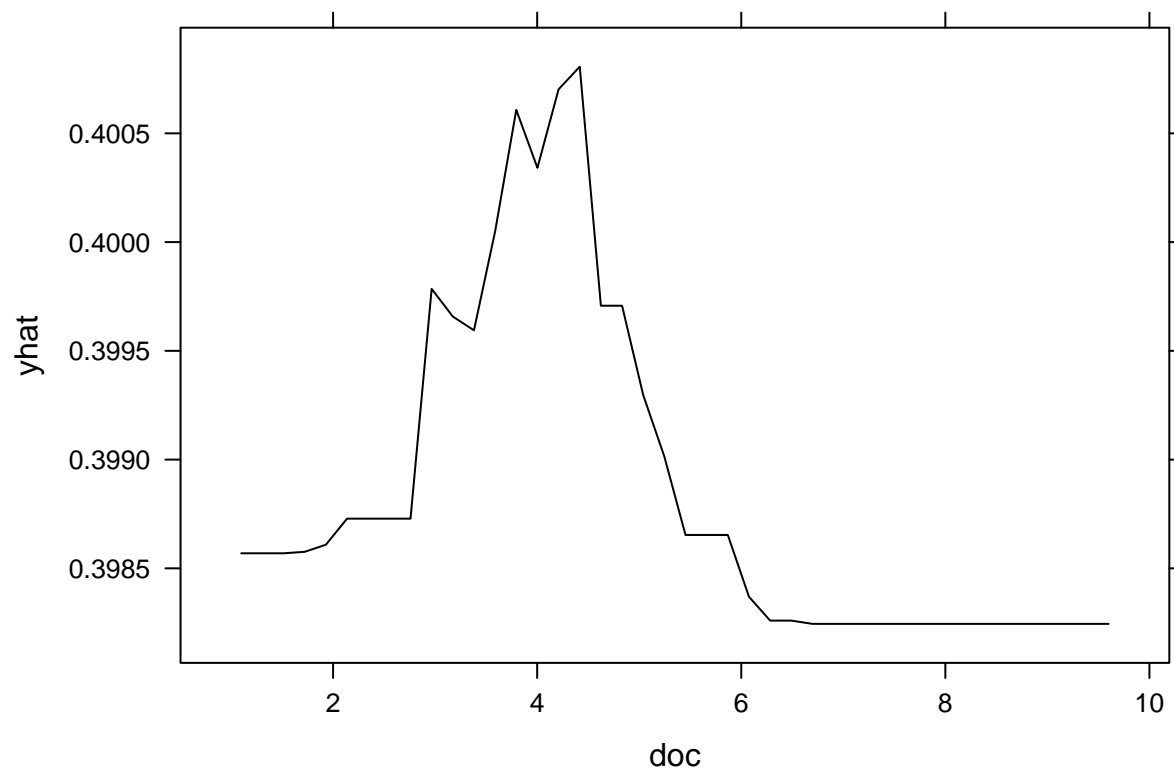
```
partial(cforest.st, pred.var="lake_area_ha", train=modvars.accndvi, type="regression", plot=T)
```



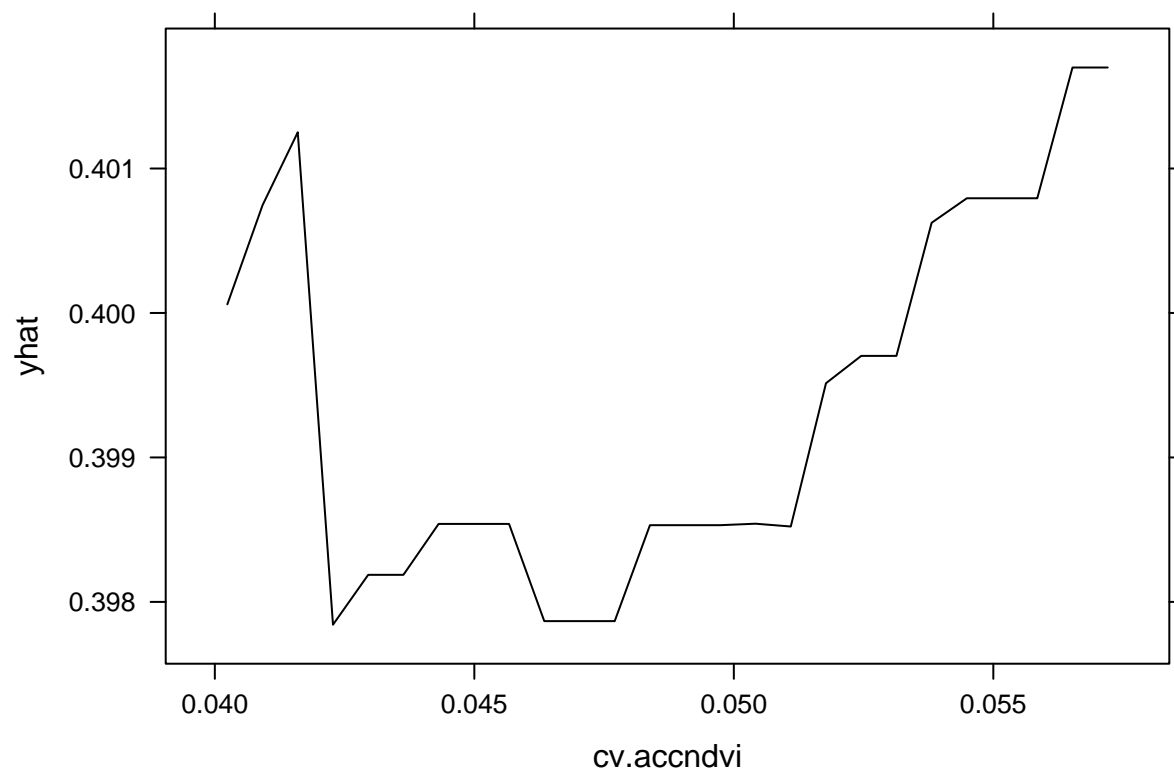
```
partial(cforest.st, pred.var="chla", train=modvars.accndvi, type="regression", plot=T)
```



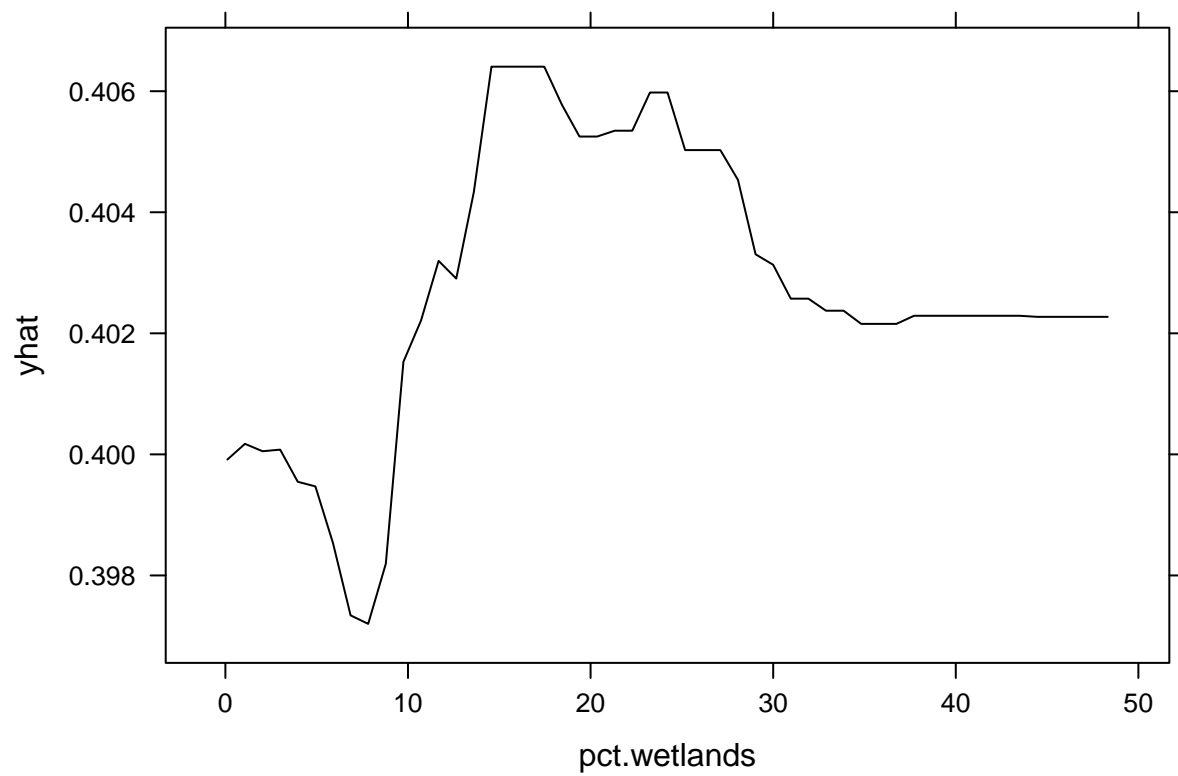
```
partial(cforest.st, pred.var="doc", train=modvars.accndvi, type="regression", plot=T)
```



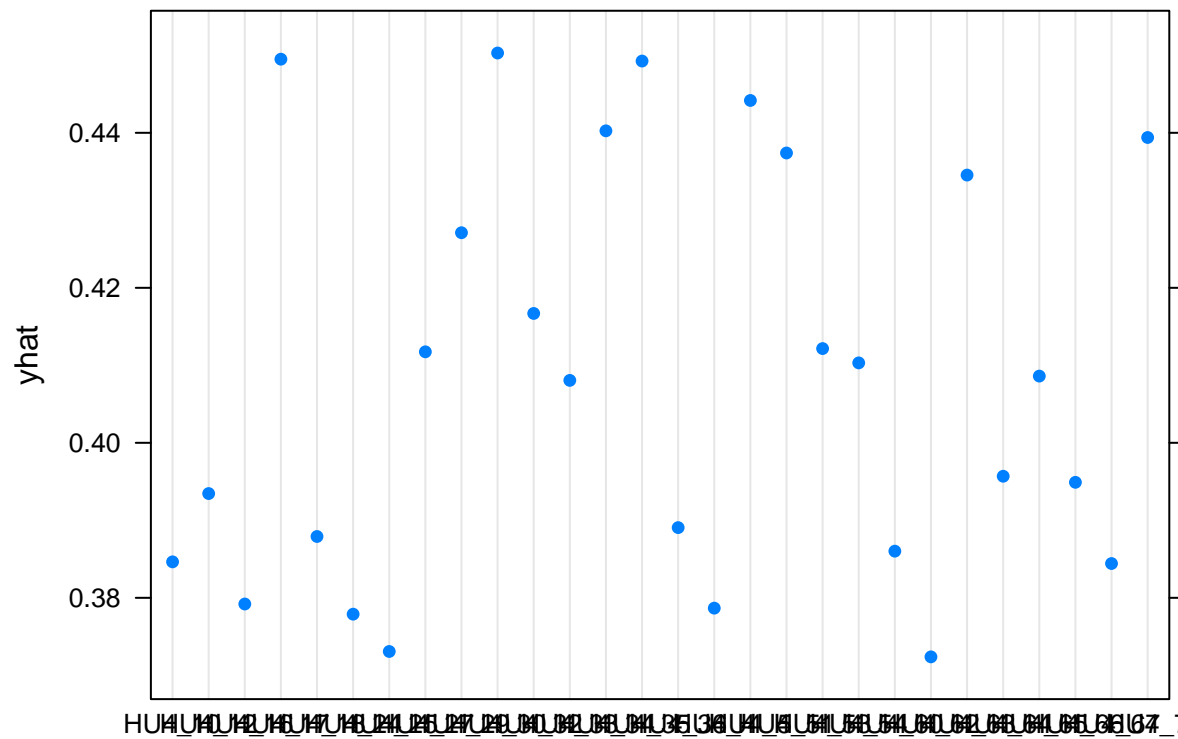
```
partial(cforest.st, pred.var="cv.accndvi", train=modvars.accndvi, type="regression", plot=T)
```



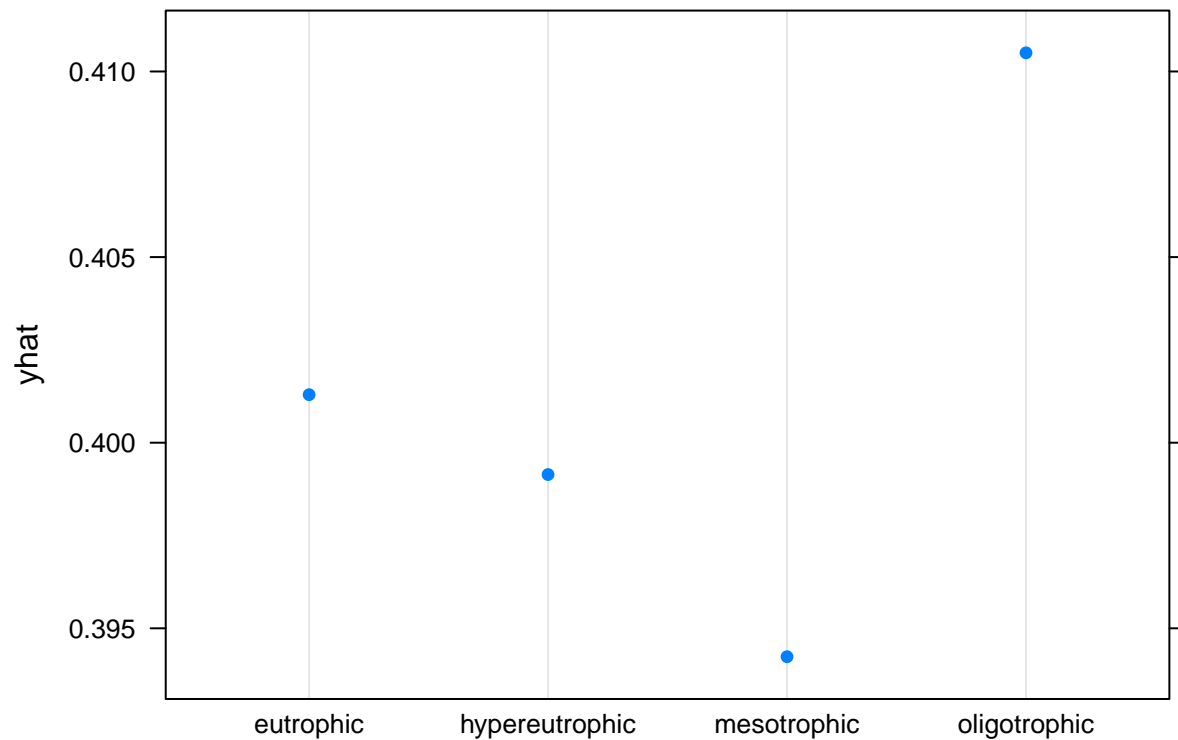
```
partial(cforest.st, pred.var="pct.wetlands", train=modvars.accndvi, type="regression", plot=T)
```



```
partial(cforest.st, pred.var="hu4_zoneid", train=modvars.accndvi, type="regression", plot=T)
```

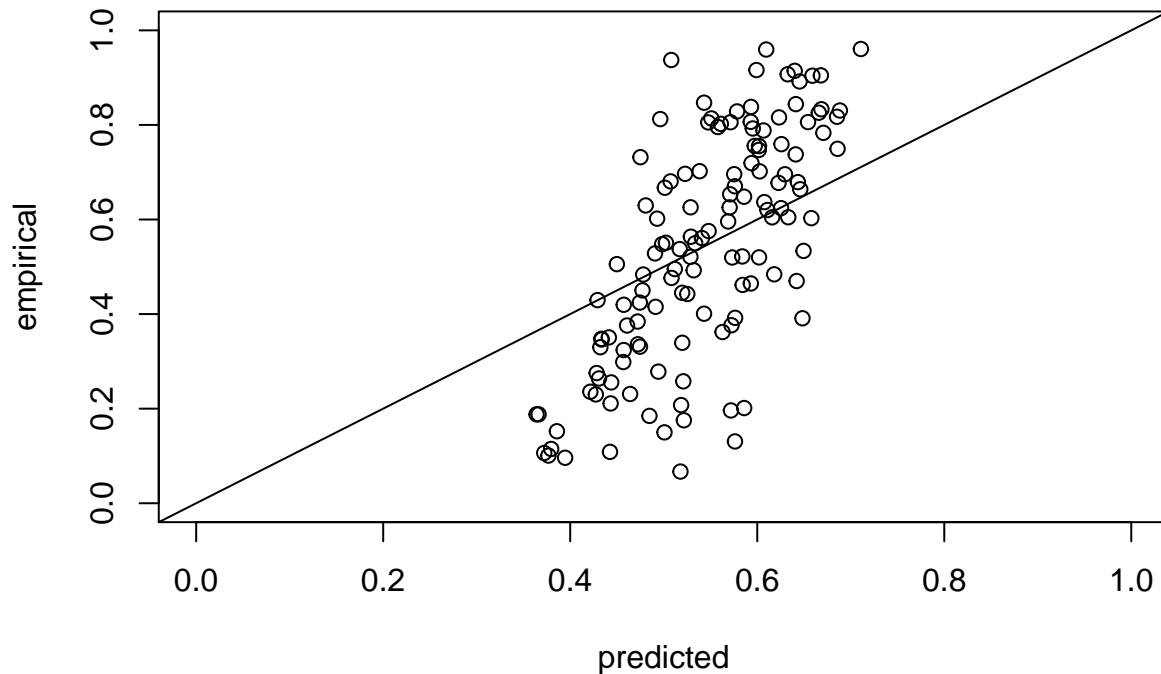


```
partial(cforest.st, pred.var="tsi.cat", train=modvars.accndvi, type="regression", plot=T)
```



```
cforest.lt<-party::cforest(accndvicoh.ts2 ~ shoredev + lake_area_ha + maxdepth + pct.ag + chla + tsi.ca
+ cv.accndvi+ pct.wetlands + doc + prcp.normal,
data=modvars.accndvi, controls=cforest_control(ntree=80000))
predcoh.lt<-predict(cforest.lt, newdata=modvars.accndvi)
# hist(predcoh.lt)
# hist(modvars.accndvi$accndvicoh.ts2)
plot(predcoh.lt, modvars.accndvi$accndvicoh.ts2, xlab="predicted", ylab="empirical", main="Coherence, l
xlim=c(0,1), ylim=c(0,1))
abline(a=0,b=1)
```

## Coherence, long st



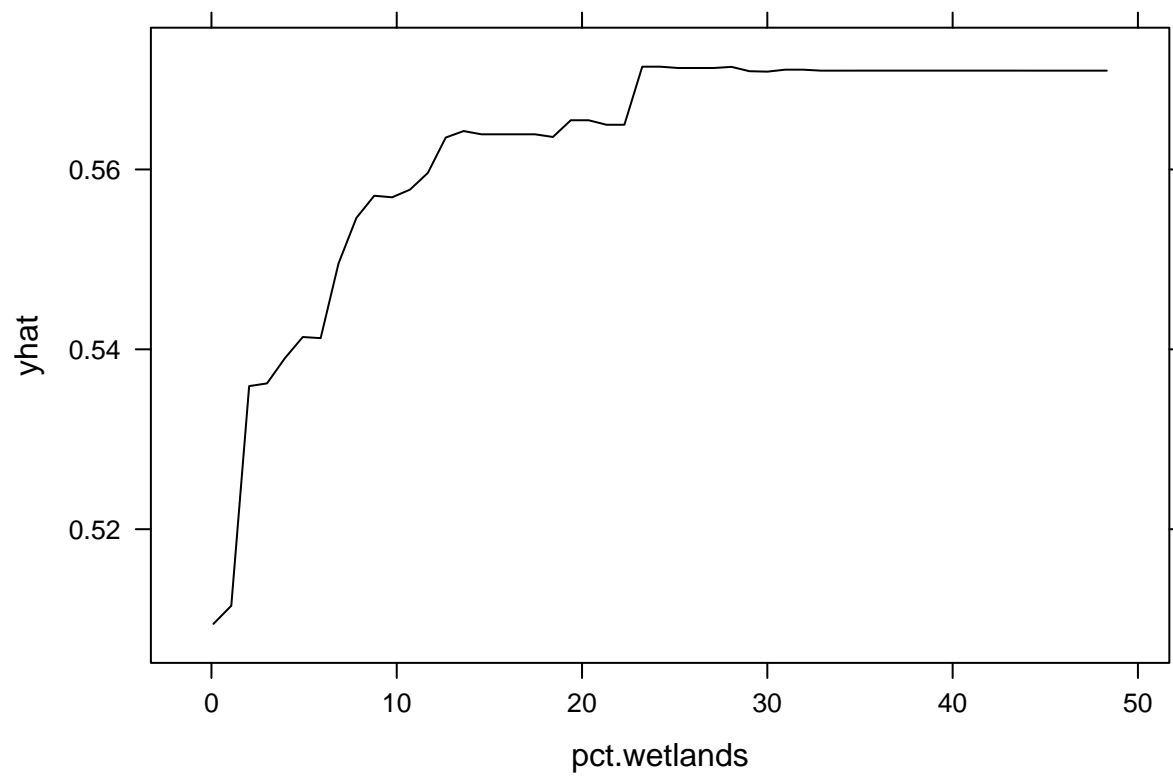
```
cor.test(predcoh.lt,modvars.accndvi$accndvicoh.ts2)
```

```
##
## Pearson's product-moment correlation
##
## data: predcoh.lt and modvars.accndvi$accndvicoh.ts2
## t = 11.478, df = 129, p-value < 2.2e-16
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## 0.6141663 0.7864509
## sample estimates:
## cor
## 0.7108112
```

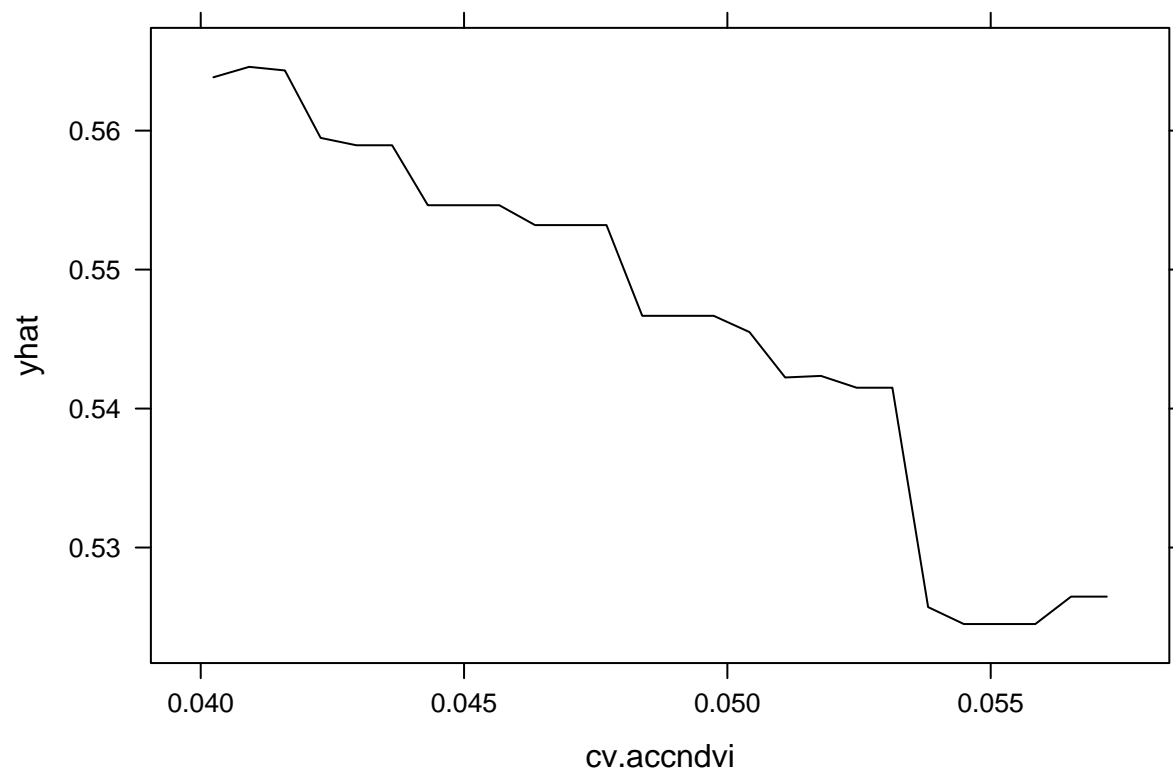
```
varimp.coh.lt<-varimp(cforest.lt)
print(varimp.coh.lt[order(varimp.coh.lt, decreasing=T)])
```

```
## pct.wetlands cv.accndvi hu4_zoneid tsi.cat doc
## 1.626927e-03 6.584396e-04 2.470309e-04 8.565307e-06 -4.165369e-05
## prcp.normal lake_area_ha chla shoredev pct.ag
## -5.131429e-05 -1.174315e-04 -1.321690e-04 -2.347191e-04 -2.685574e-04
## maxdepth
## -4.483252e-04
```

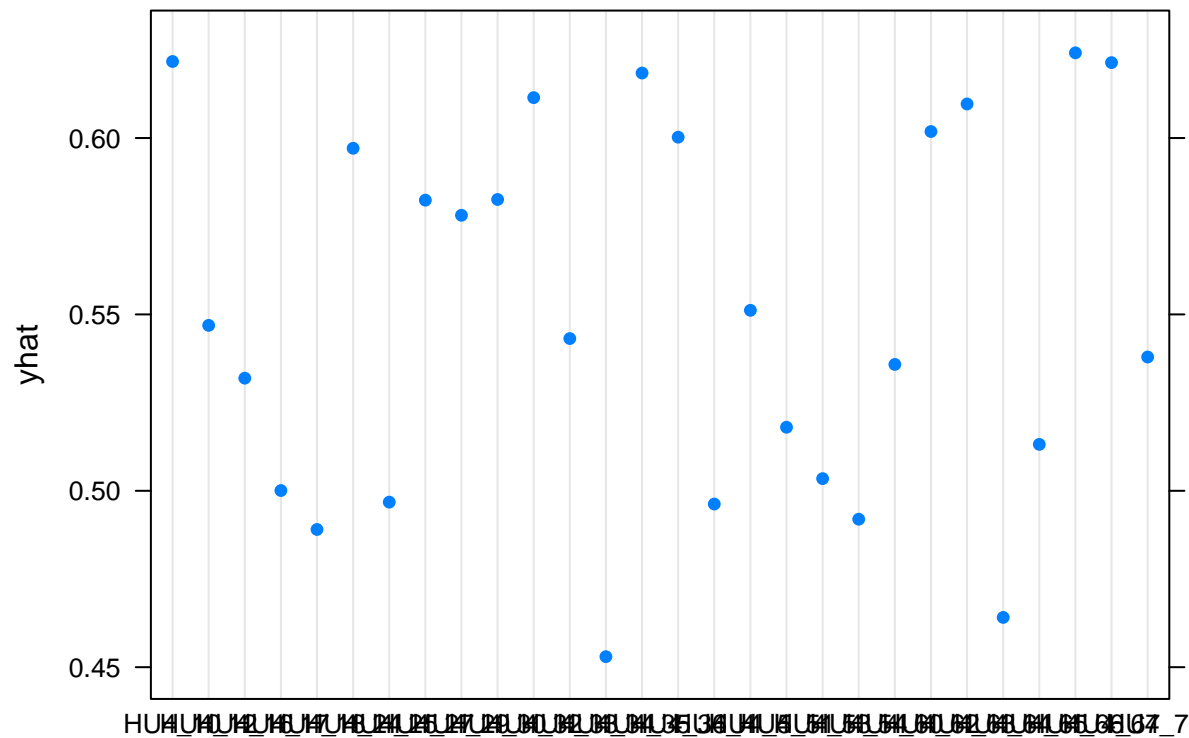
```
partial(cforest.lt, pred.var="pct.wetlands", train=modvars.accndvi, type="regression", plot=T)
```



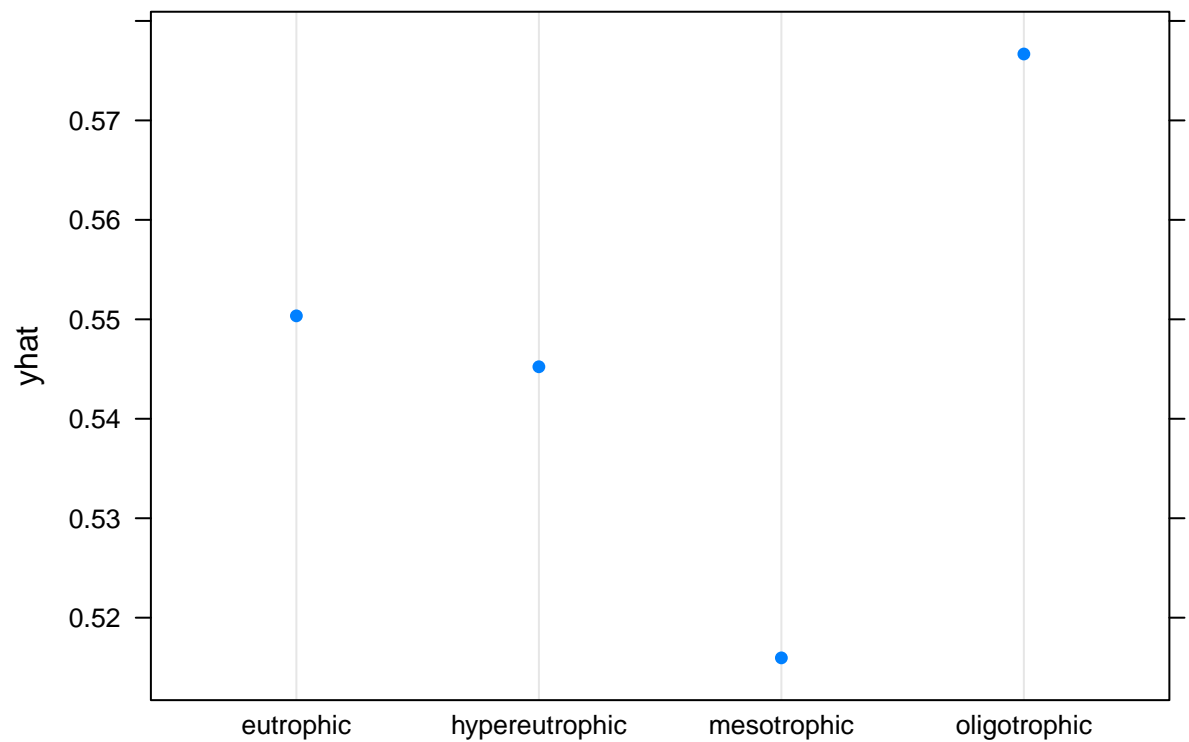
```
partial(cforest.lt, pred.var="cv.accndvi", train=modvars.accndvi, type="regression", plot=T)
```



```
partial(cforest.lt, pred.var="hu4_zoneid", train=modvars.accndvi, type="regression", plot=T)
```

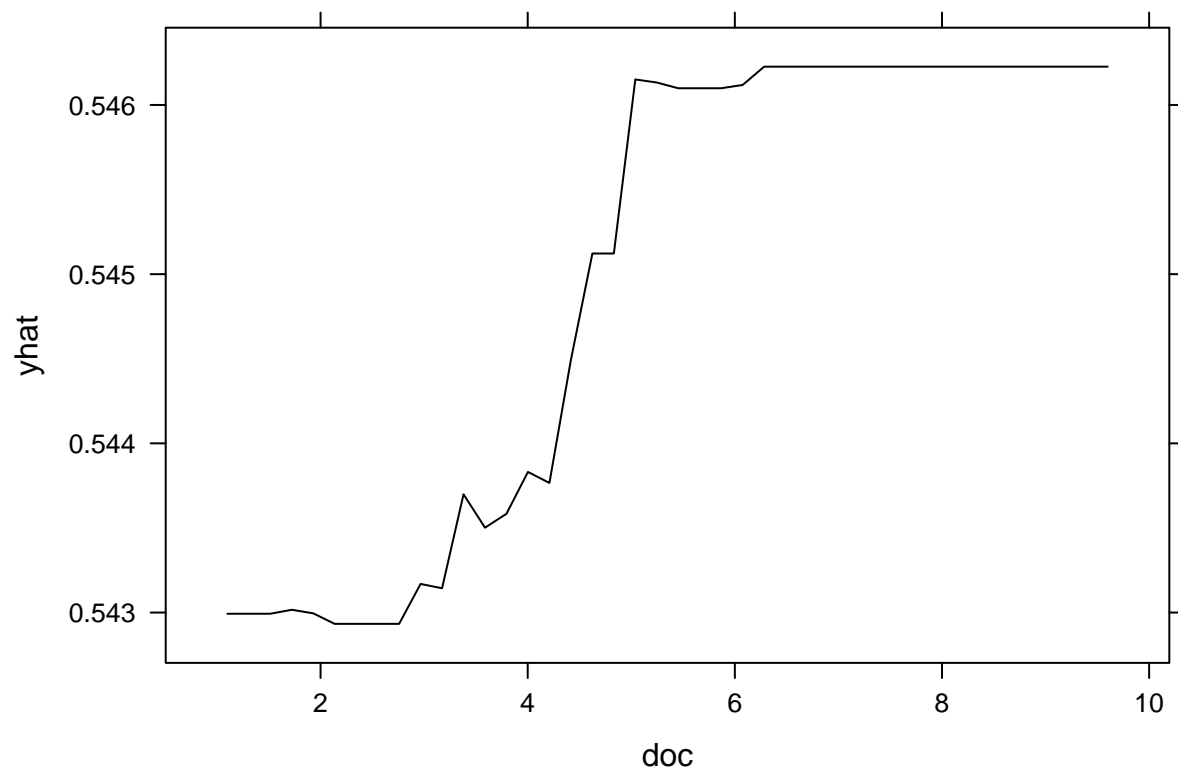


```
partial(cforest.lt, pred.var="tsi.cat", train=modvars.accndvi, type="regression", plot=T)
```

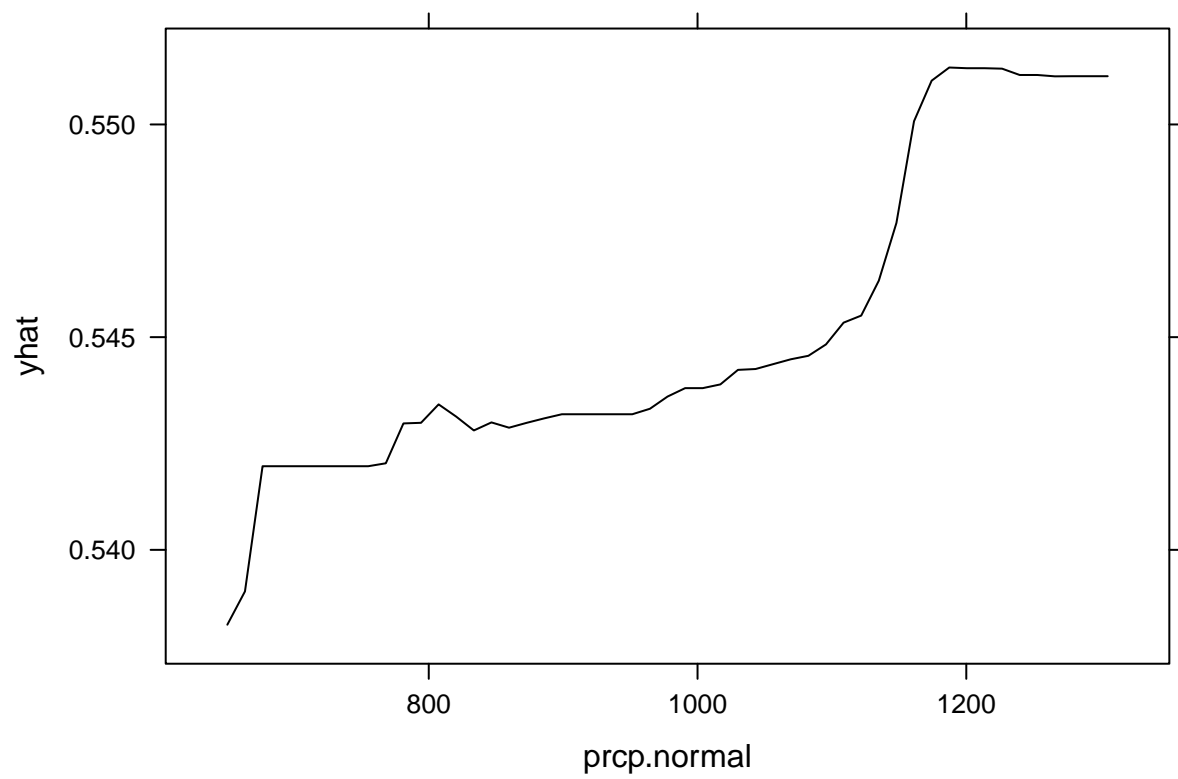


```
partial(cforest.lt, pred.var="doc", train=modvars.accndvi, type="regression", plot=T)
```

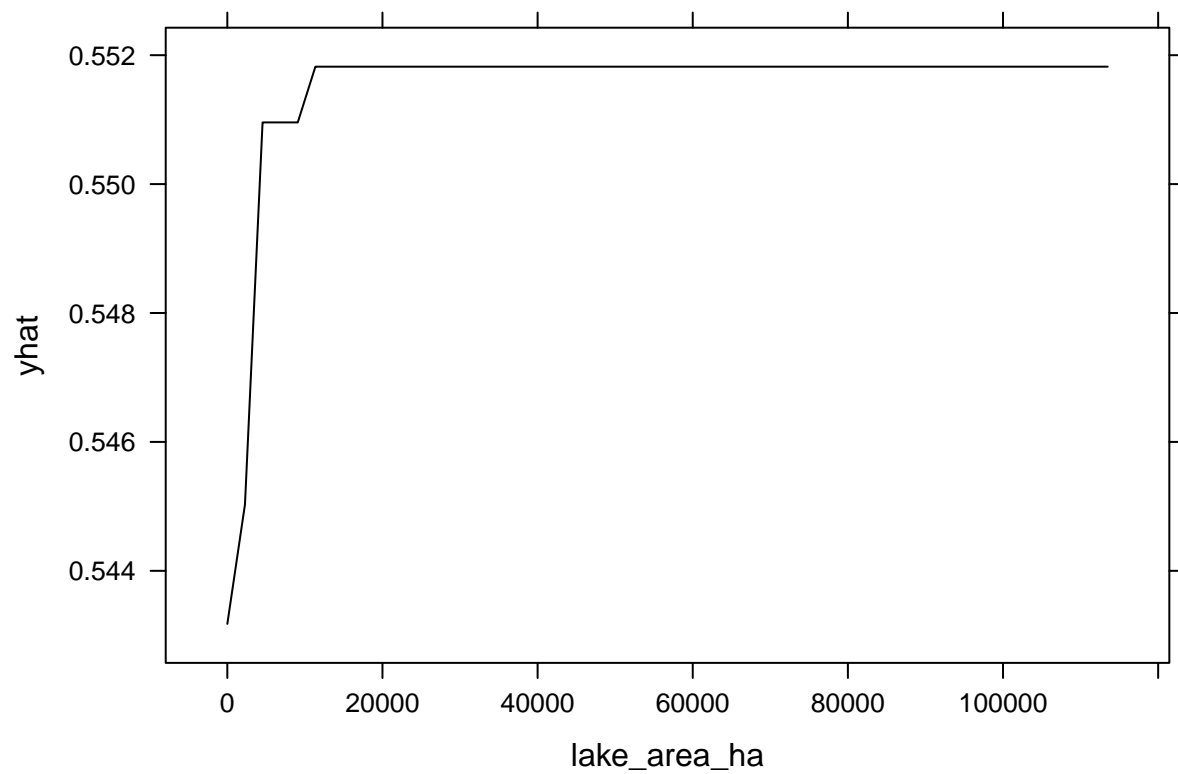




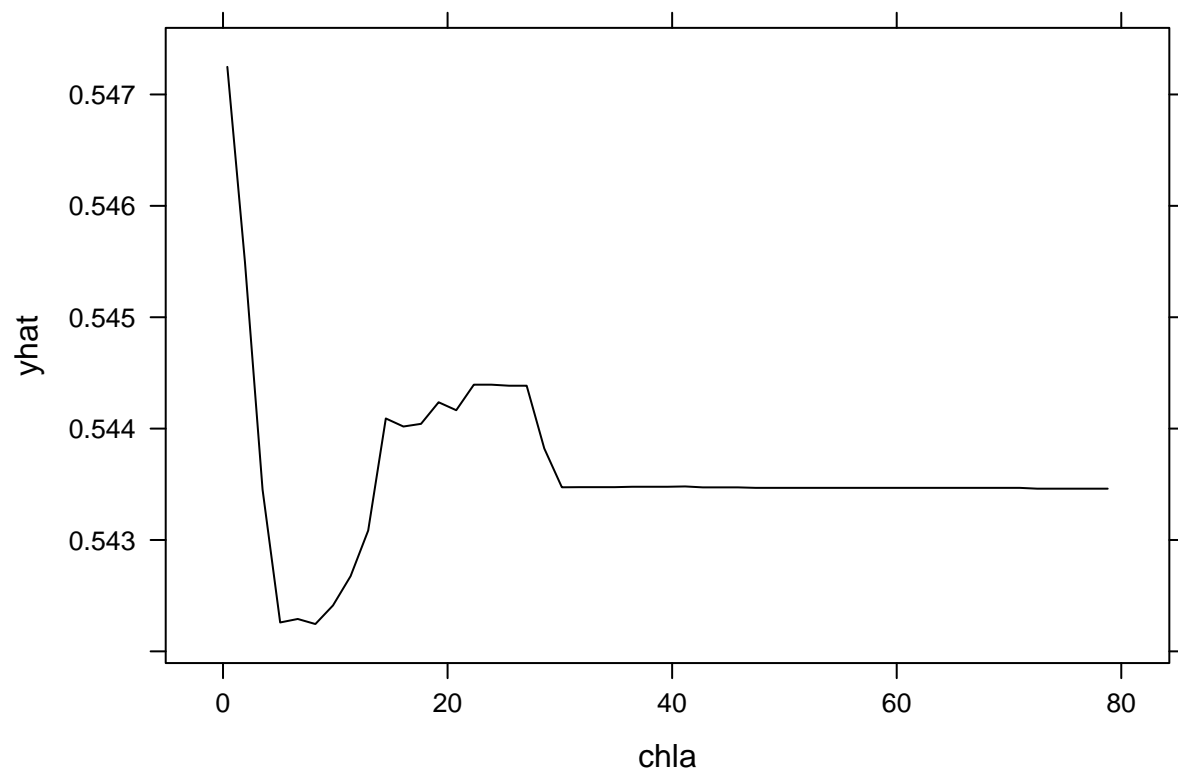
```
partial(cforest.lt, pred.var="prcp.normal", train=modvars.accndvi, type="regression", plot=T)
```



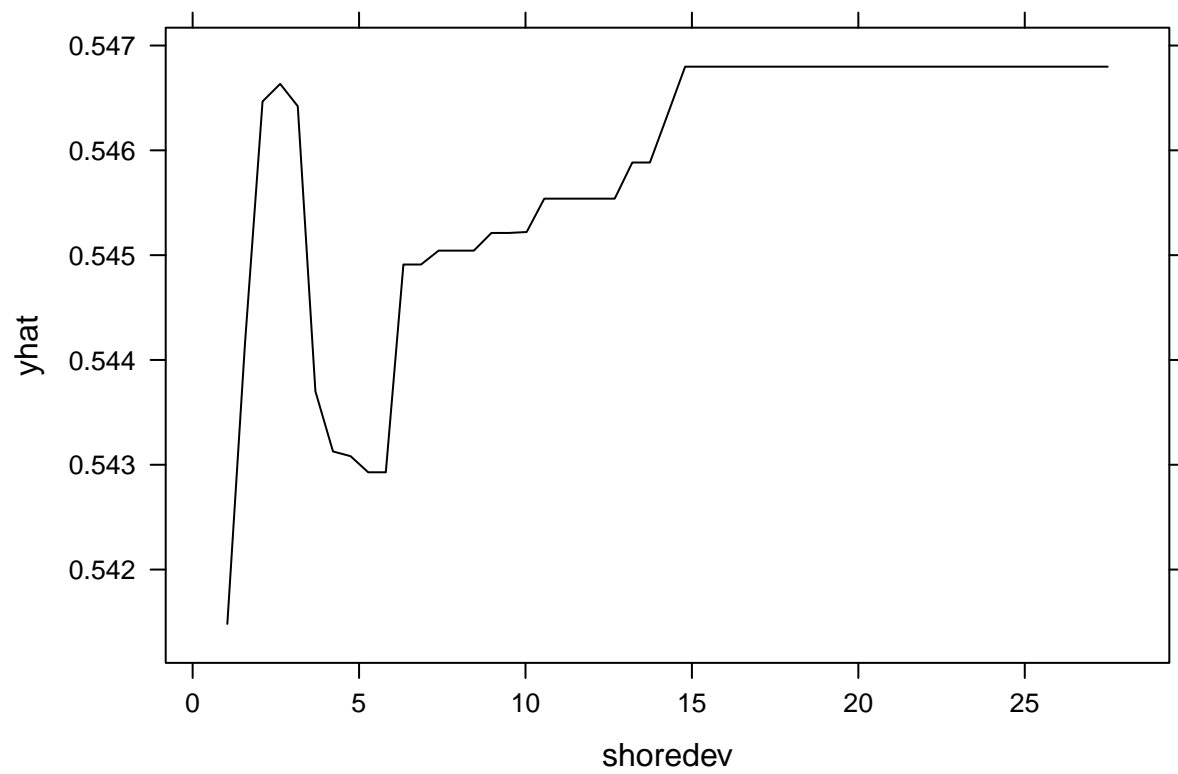
```
partial(cforest.lt, pred.var="lake_area_ha", train=modvars.accndvi, type="regression", plot=T)
```



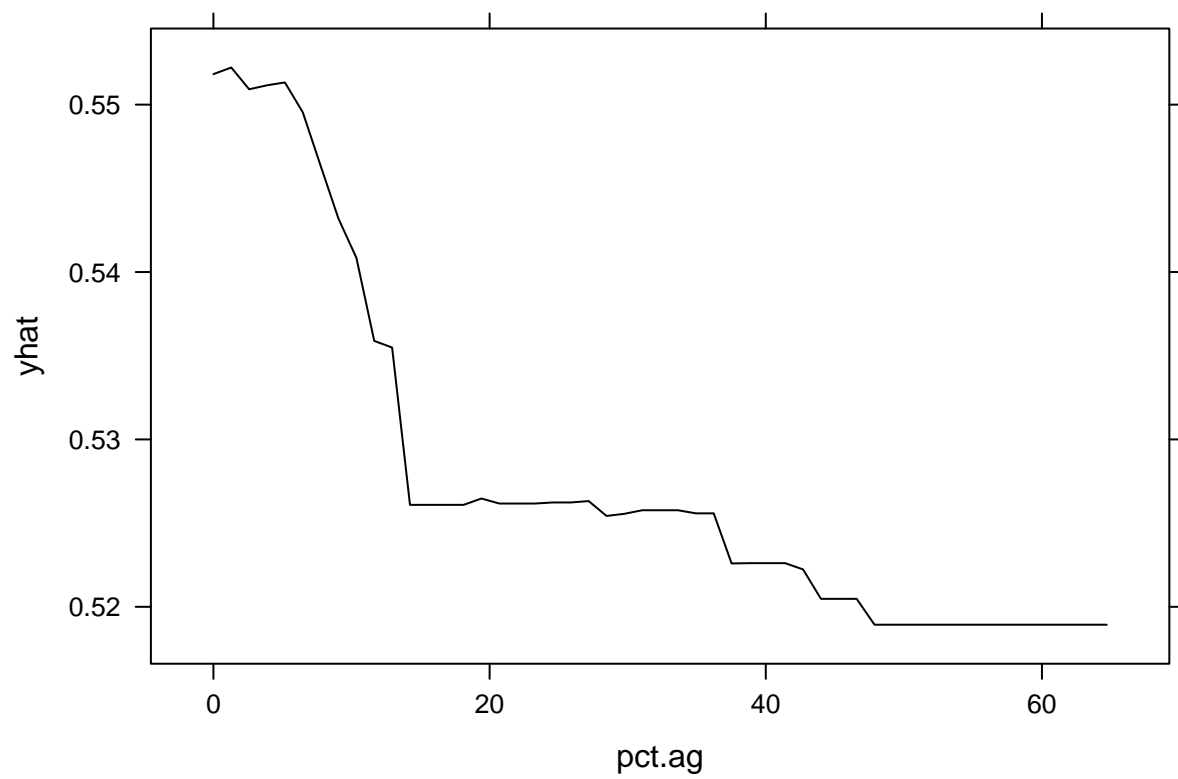
```
partial(cforest.lt, pred.var="chla", train=modvars.accndvi, type="regression", plot=T)
```



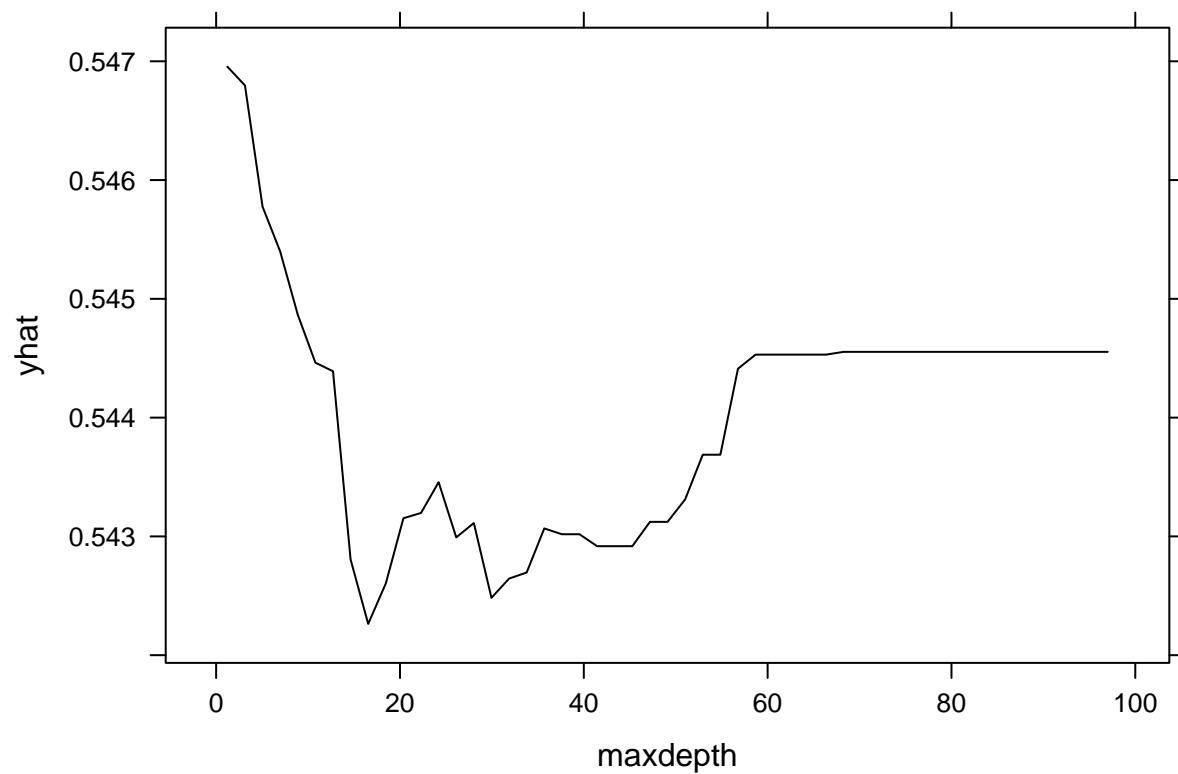
```
partial(cforest.lt, pred.var="shoredew", train=modvars.accndvi, type="regression", plot=T)
```



```
partial(cforest.lt, pred.var="pct.ag", train=modvars.acndvi, type="regression", plot=T)
```

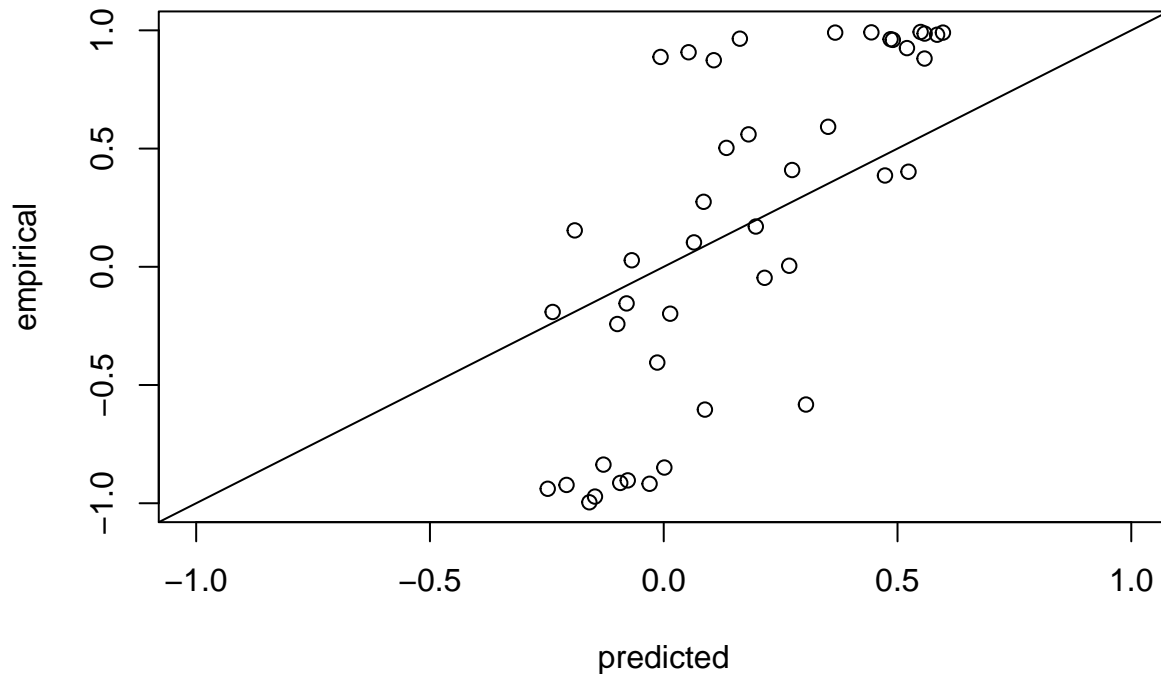


```
partial(cforest.lt, pred.var="maxdepth", train=modvars.acndvi, type="regression", plot=T)
```



```
cforest.phi.st<-cforest(cos(accndviphi.ts1) ~ shoredev + lake_area_ha + maxdepth + pct.ag + chla + tsi.
                        hu4_zoneid + cv.accndvi+ pct.wetlands + doc + prcp.normal,
                        data=modvars.accndvi.phist,controls=cforest_control(ntree=40000))
predphi.st<-predict(cforest.phi.st, newdata=modvars.accndvi.phist)
# hist(predphi.st)
# hist(cos(modvars.accndvi.phist$accndviphi.ts1))
plot(predphi.st, cos(modvars.accndvi.phist$accndviphi.ts1), xlab="predicted", ylab="empirical", main="c
      xlim=c(-1,1), ylim=c(-1,1))
abline(a=0,b=1)
```

## cos(phase), short ts



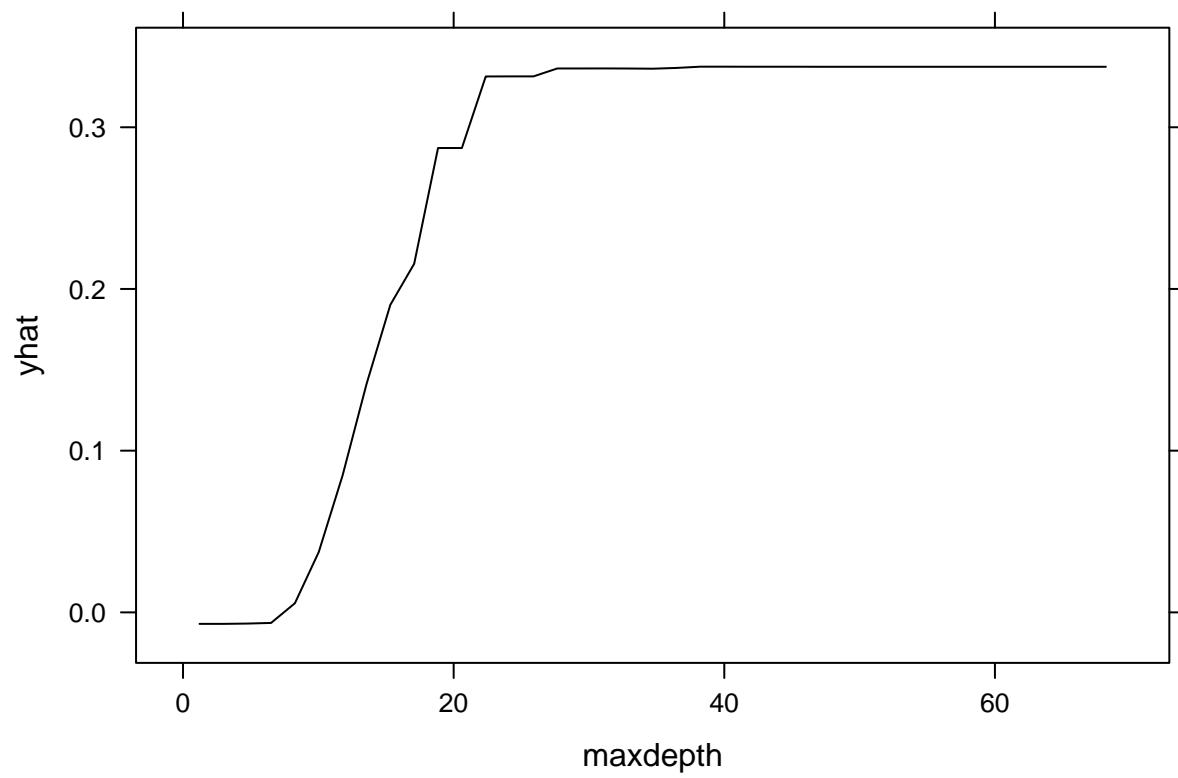
```
cor.test(predphi.st,cos(modvars.accndvi.phist$accndviphi.ts1))
```

```
##
## Pearson's product-moment correlation
##
## data: predphi.st and cos(modvars.accndvi.phist$accndviphi.ts1)
## t = 7.2976, df = 41, p-value = 6.285e-09
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.5829267 0.8582544
## sample estimates:
##      cor
## 0.7516695
```

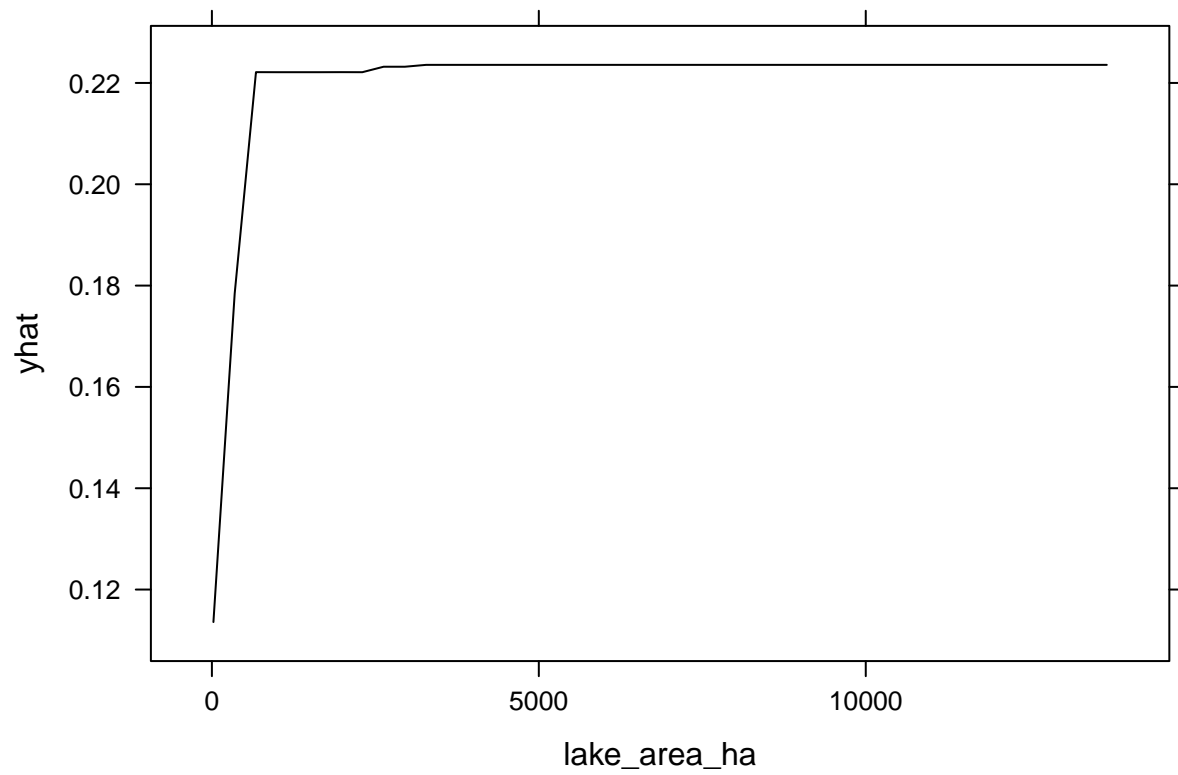
```
varimp.phi.st<-varimp(cforest.phi.st)
print(varimp.phi.st[order(varimp.phi.st,decreasing=TRUE)])
```

```
##      maxdepth lake_area_ha      doc  cv.accndvi  shoredev
## 6.759071e-02 1.905140e-02 3.437781e-06 -1.745340e-04 -4.132123e-04
## pct.wetlands prcp.normal      chla      pct.ag      tsi.cat
## -7.192697e-04 -3.650594e-03 -4.078876e-03 -5.249392e-03 -8.743358e-03
## hu4_zoneid
## -2.458269e-02
```

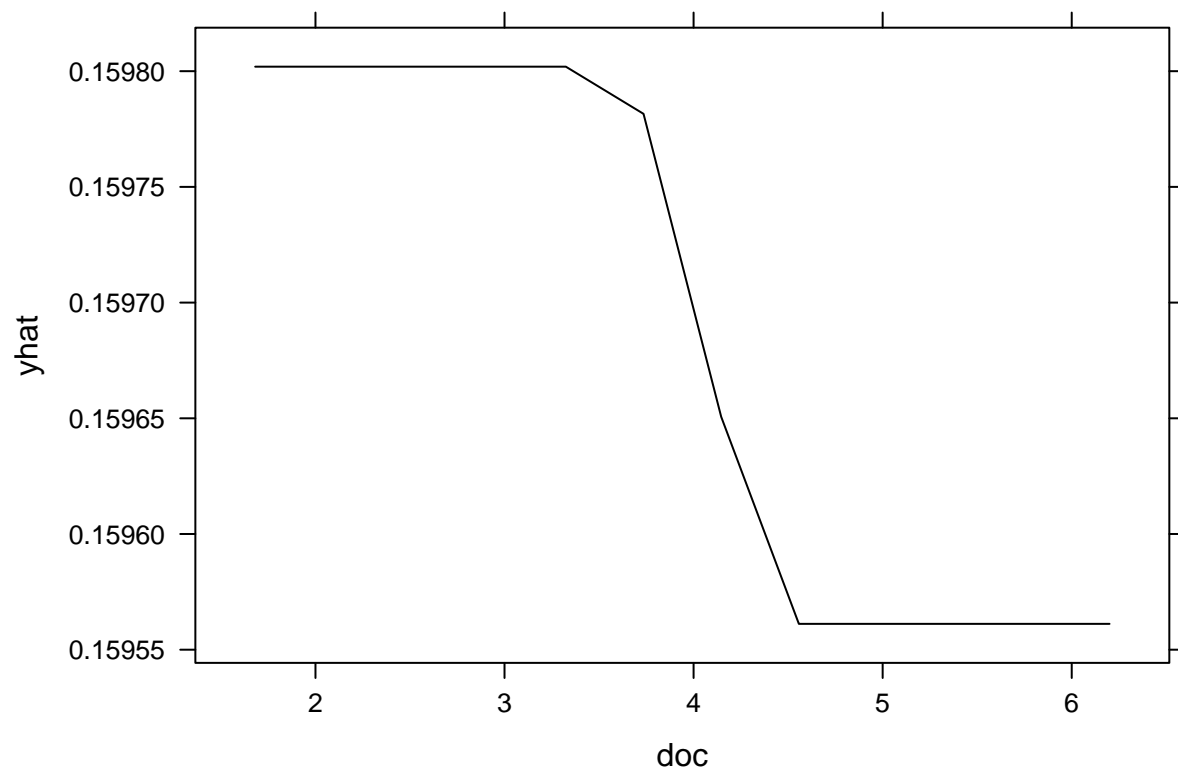
```
partial(cforest.phi.st, pred.var="maxdepth", train=modvars.accndvi.phist, type="regression", plot=T)
```



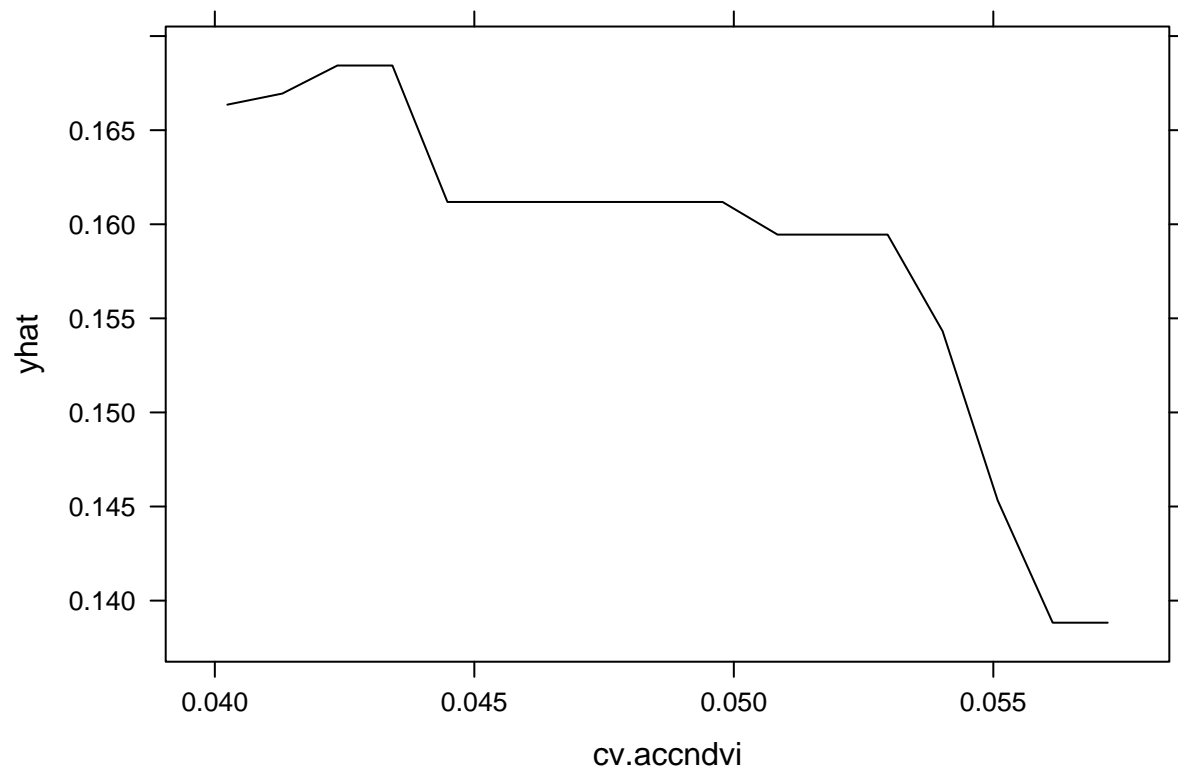
```
partial(cforest.phi.st, pred.var="lake_area_ha", train=modvars.accndvi.phist, type="regression", plot=T)
```



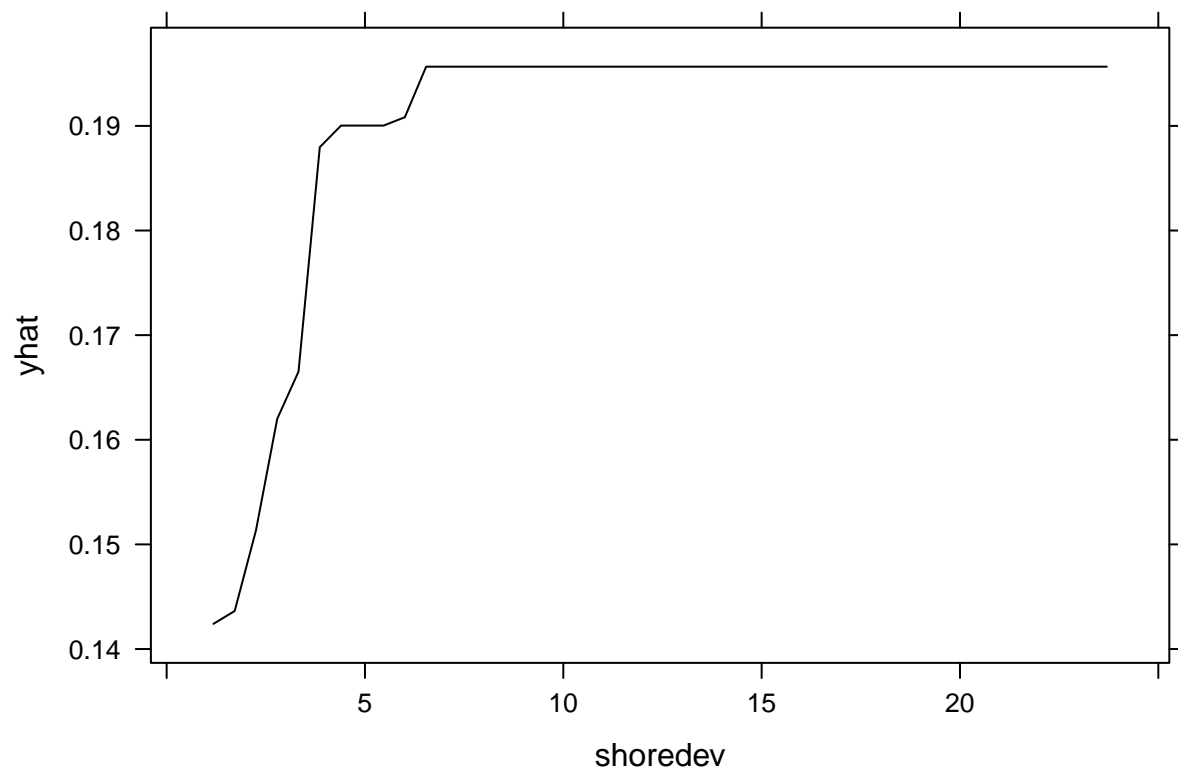
```
partial(cforest.phi.st, pred.var="doc", train=modvars.accndvi.phist, type="regression", plot=T)
```



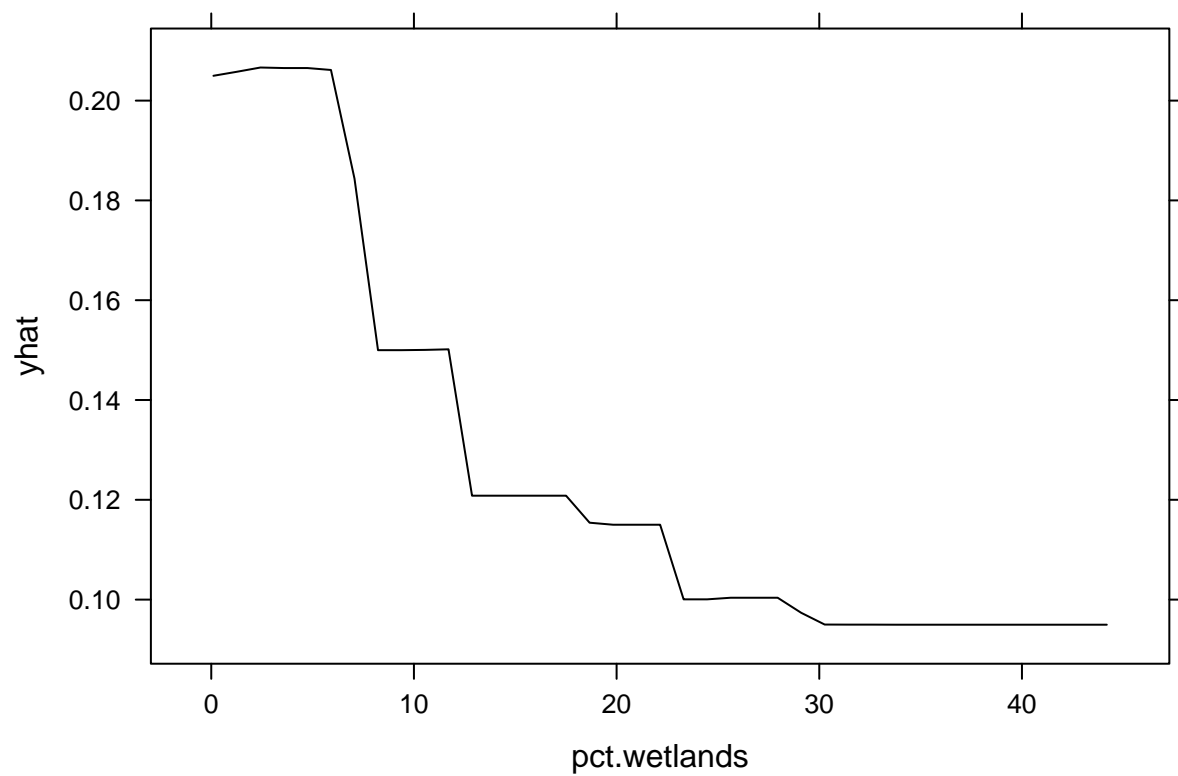
```
partial(cforest.phi.st, pred.var="cv.accndvi", train=modvars.accndvi.phist, type="regression", plot=T)
```



```
partial(cforest.phi.st, pred.var="shoreddev", train=modvars.accndvi.phist, type="regression", plot=T)
```

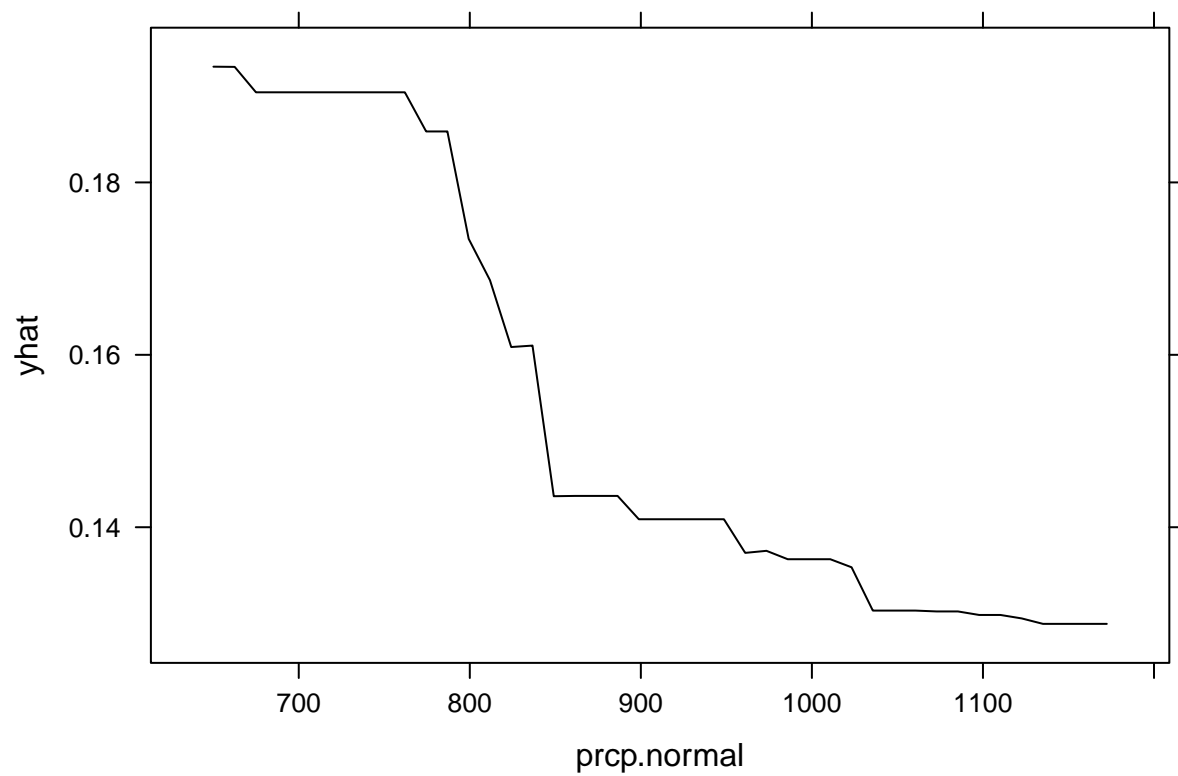


```
partial(cforest.phi.st, pred.var="pct.wetlands", train=modvars.acndvi.phist, type="regression", plot=T)
```

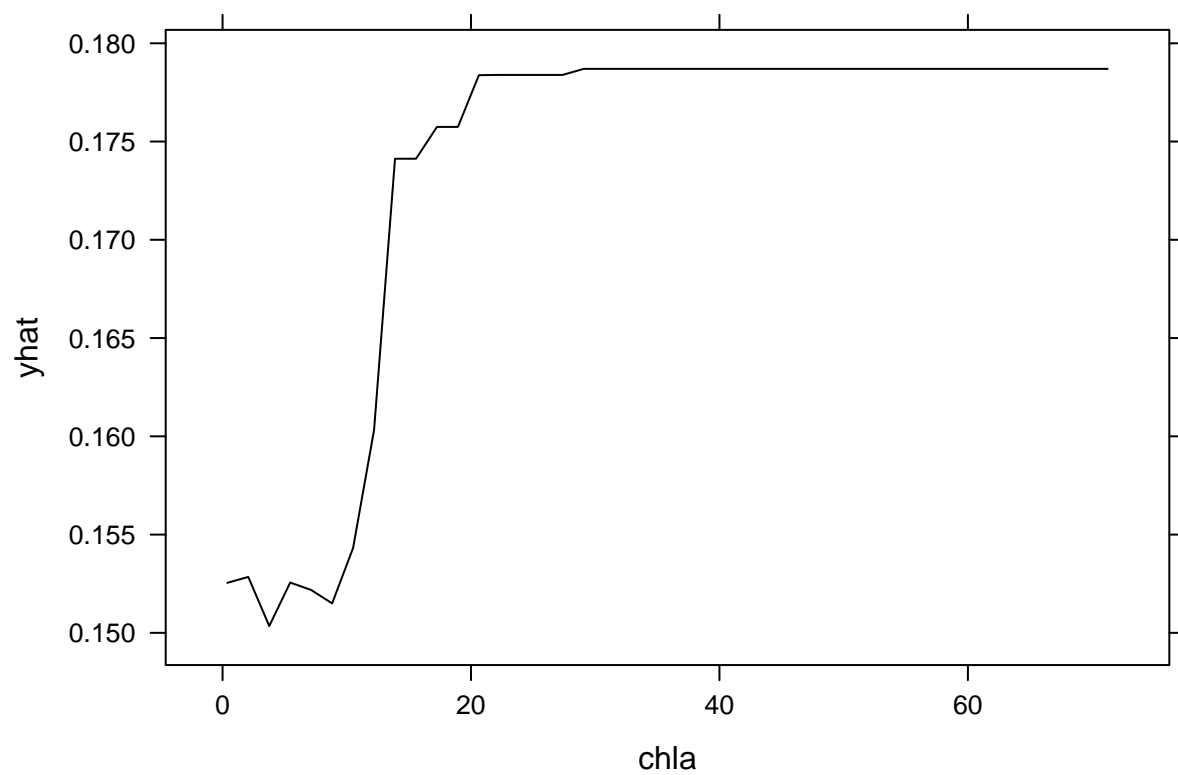


```
partial(cforest.phi.st, pred.var="prcp.normal", train=modvars.acndvi.phist, type="regression", plot=T)
```

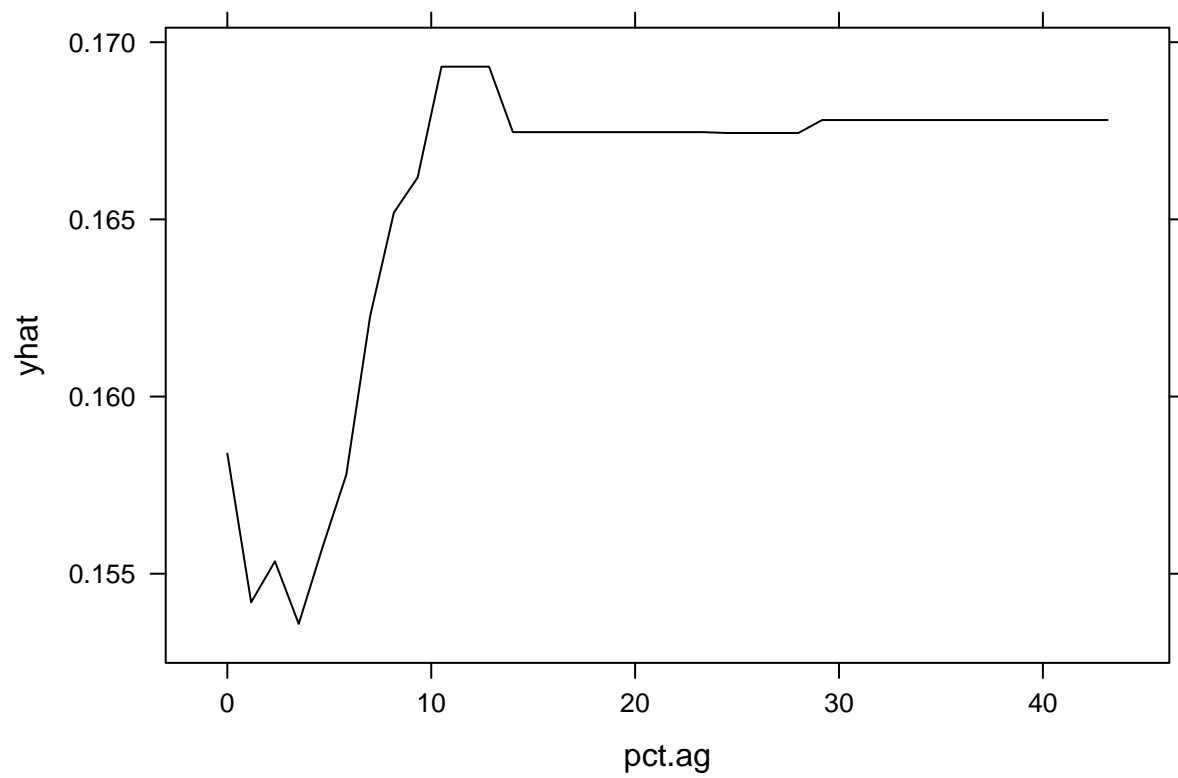




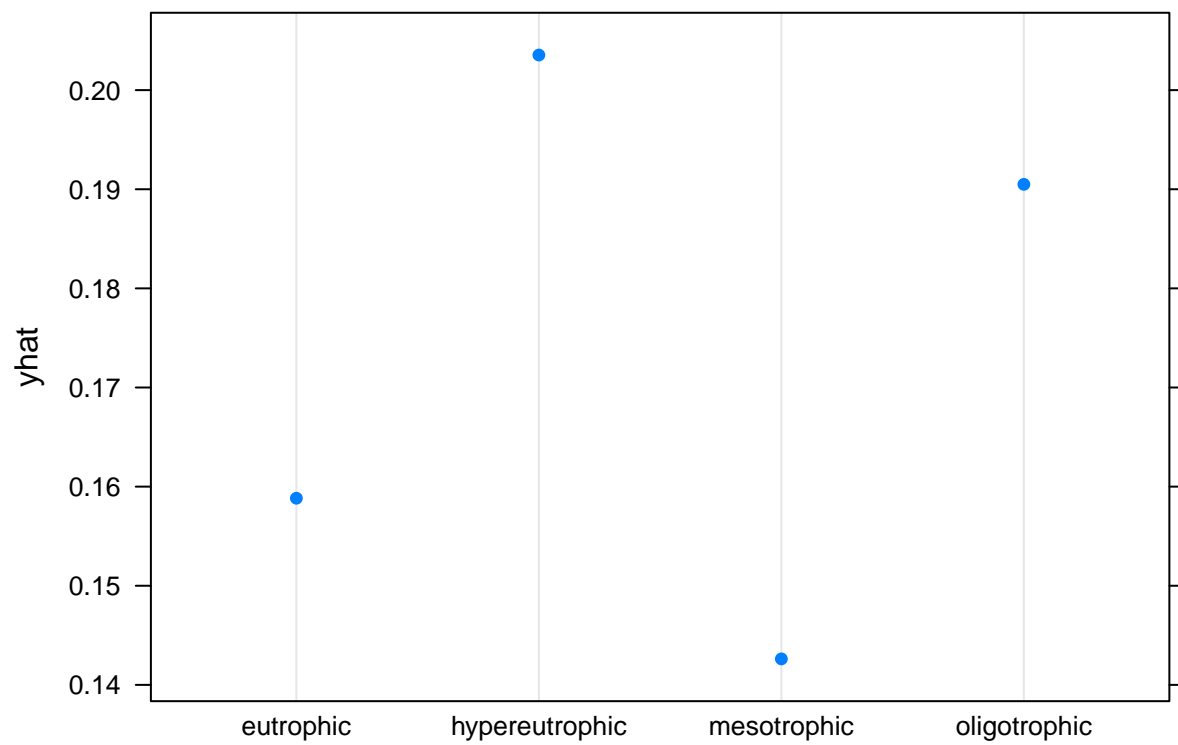
```
partial(cforest.phi.st, pred.var="chla", train=modvars.accndvi.phist, type="regression", plot=T)
```



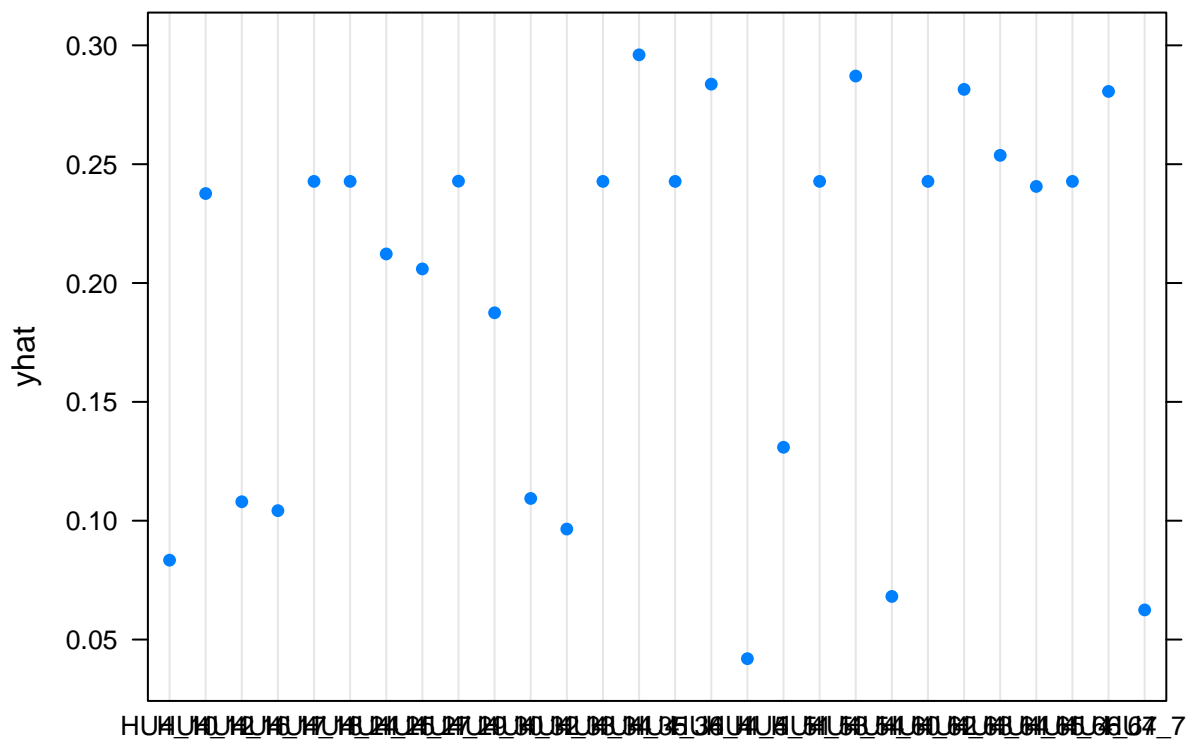
```
partial(cforest.phi.st, pred.var="pct.ag", train=modvars.accndvi.phist, type="regression", plot=T)
```



```
partial(cforest.phi.st, pred.var="tsi.cat", train=modvars.accndvi.phist, type="regression", plot=T)
```

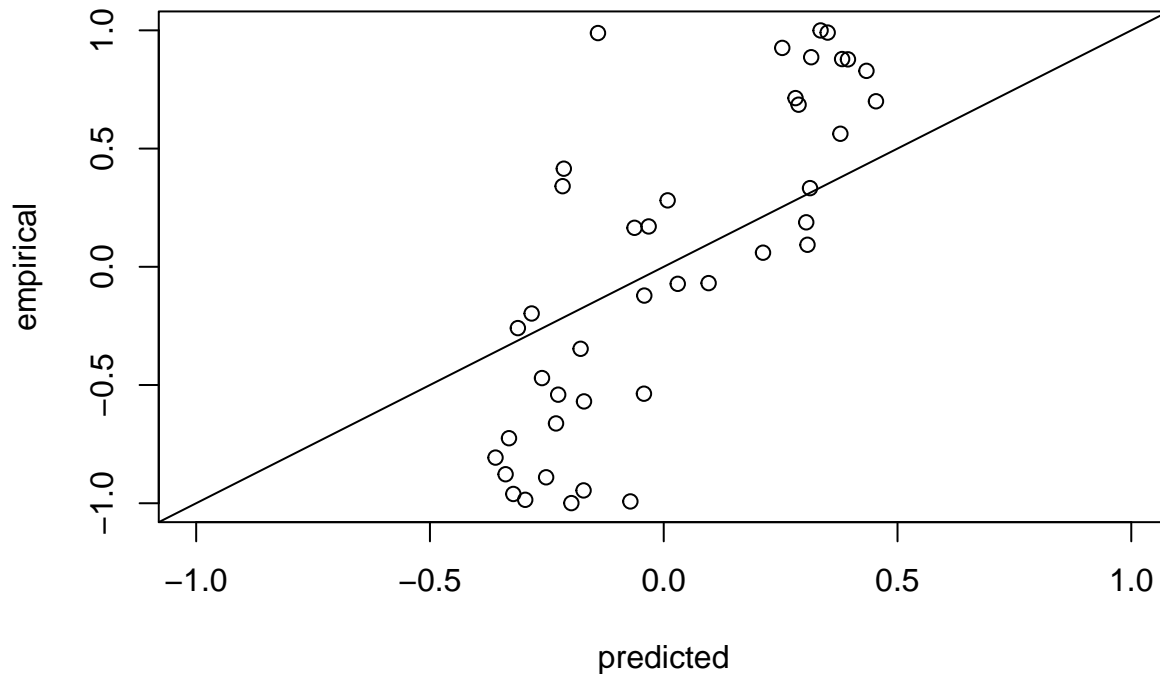


```
partial(cforest.phi.st, pred.var="hu4_zoneid", train=modvars.accndvi.phist, type="regression", plot=T)
```



```
cforest.phi.lt<-cforest(cos(accndviphi.ts2) ~ shoredev + lake_area_ha + maxdepth + pct.ag + chla + tsi.
                        cv.accndvi + pct.wetlands + doc + prcp.normal,
                        data=modvars.accndvi.philt,controls=cforest_control(ntree=40000))
predphi.lt<-predict(cforest.phi.lt, newdata=modvars.accndvi.philt)
# hist(predphi.lt)
# hist(cos(modvars.accndvi.philt$accndviphi.ts2))
plot(predphi.lt, cos(modvars.accndvi.philt$accndviphi.ts2), xlab="predicted", ylab="empirical", main="c
      xlim=c(-1,1), ylim=c(-1,1))
abline(a=0,b=1)
```

## cos(phase), short ts



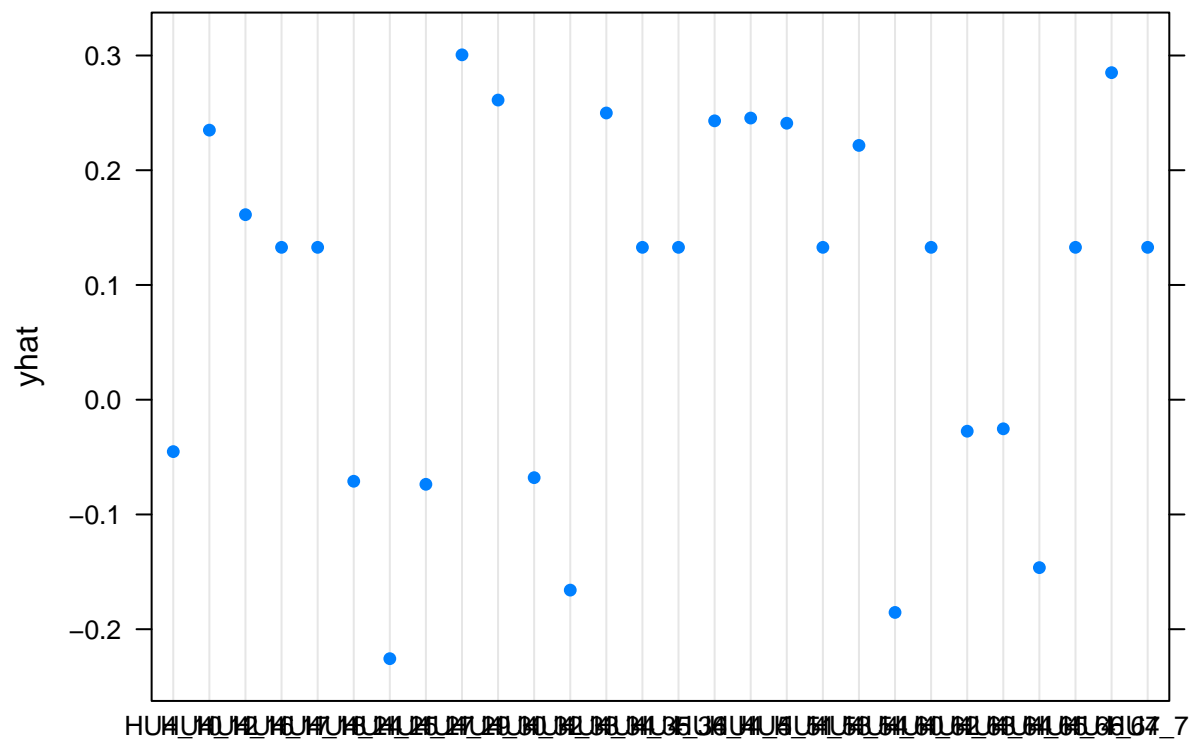
```
cor.test(predphi.lt,cos(modvars.accndvi.philt$accndviphi.ts2))
```

```
##
## Pearson's product-moment correlation
##
## data: predphi.lt and cos(modvars.accndvi.philt$accndviphi.ts2)
## t = 8.0825, df = 39, p-value = 7.35e-10
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.6392940 0.8838018
## sample estimates:
##      cor
## 0.7913103
```

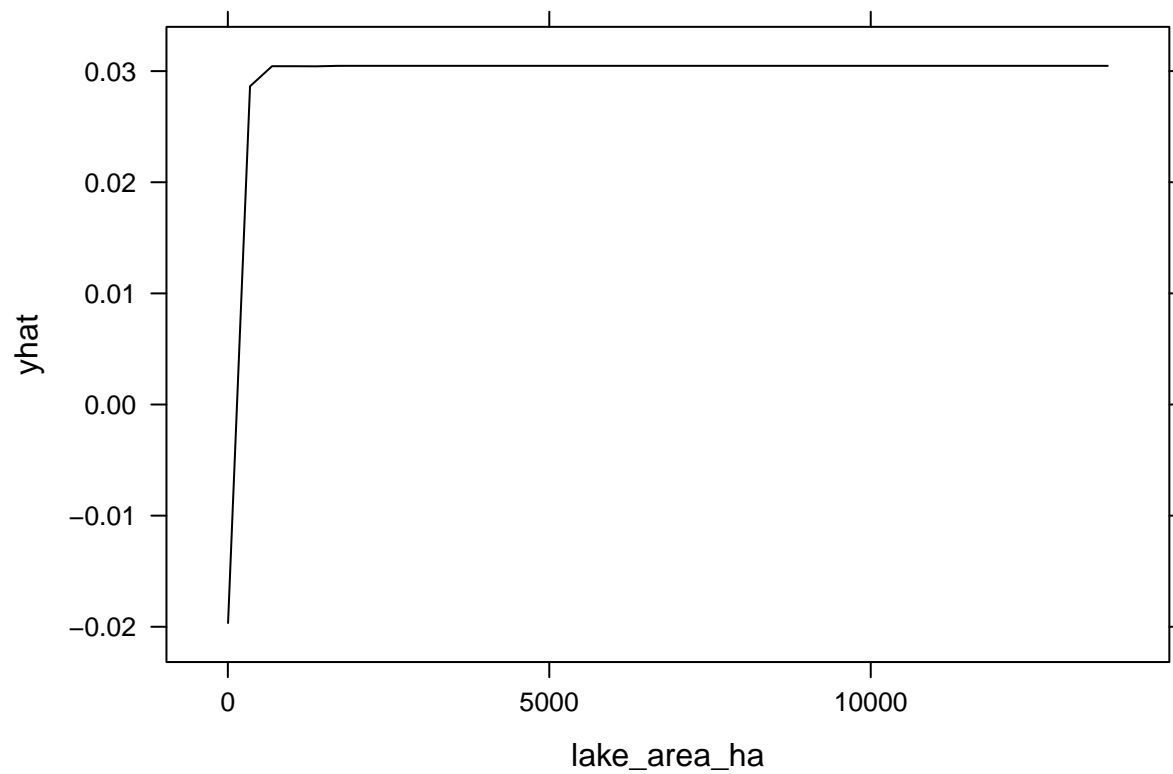
```
varimp.phi.lt<-varimp(cforest.phi.lt)
print(varimp.phi.lt[order(varimp.phi.lt, decreasing=TRUE)])
```

```
## hu4_zoneid lake_area_ha prcp.normal pct.wetlands doc
## 9.909794e-02 7.906764e-03 4.062440e-03 2.138175e-03 -3.043314e-05
## shoredev chla cv.accndvi maxdepth pct.ag
## -7.503947e-04 -3.255348e-03 -3.872061e-03 -4.251249e-03 -5.888400e-03
## tsi.cat
## -8.260009e-03
```

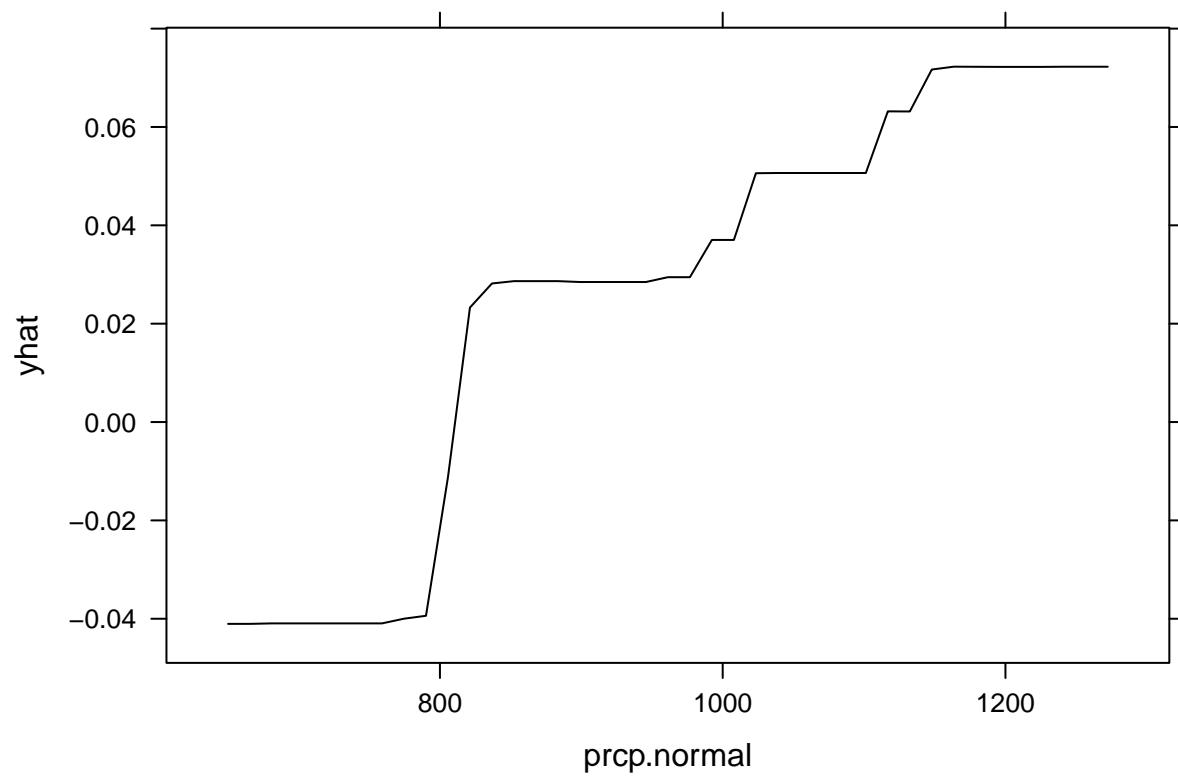
```
partial(cforest.phi.lt, pred.var="hu4_zoneid", train=modvars.accndvi.philt, type="regression", plot=T)
```



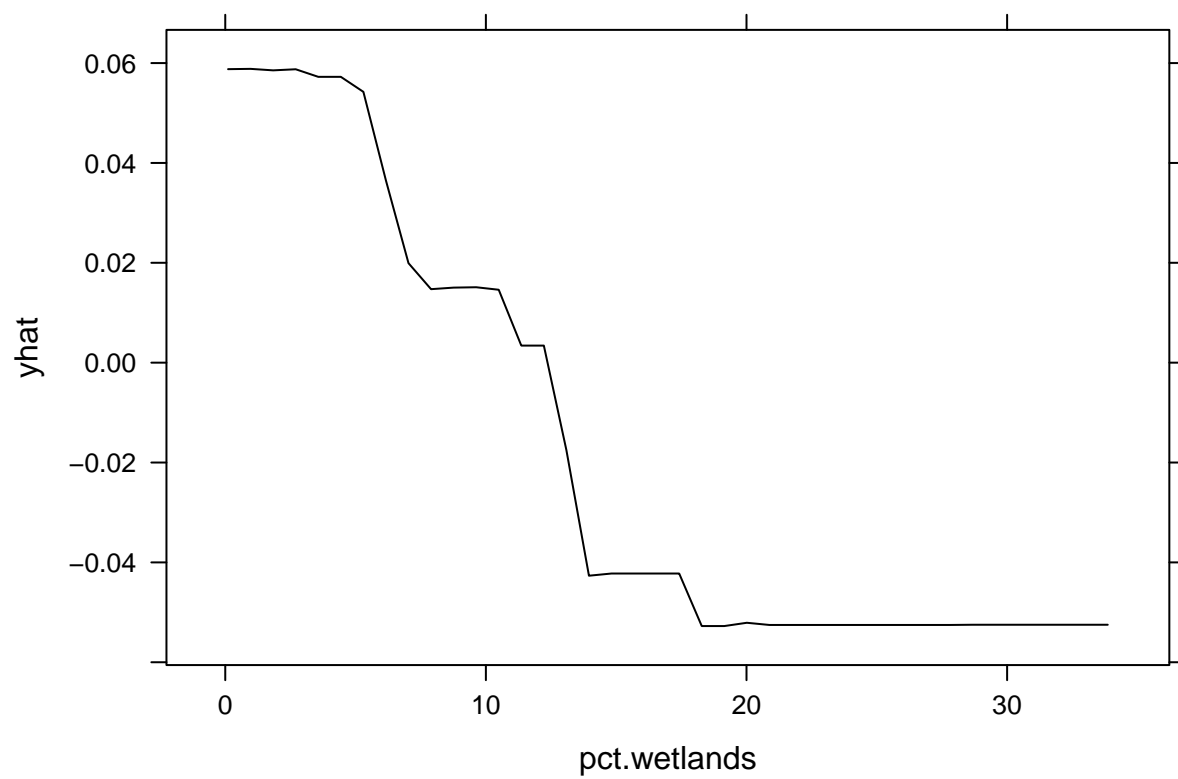
```
partial(cforest.phi.lt, pred.var="lake_area_ha", train=modvars.accndvi.philt, type="regression", plot=T)
```



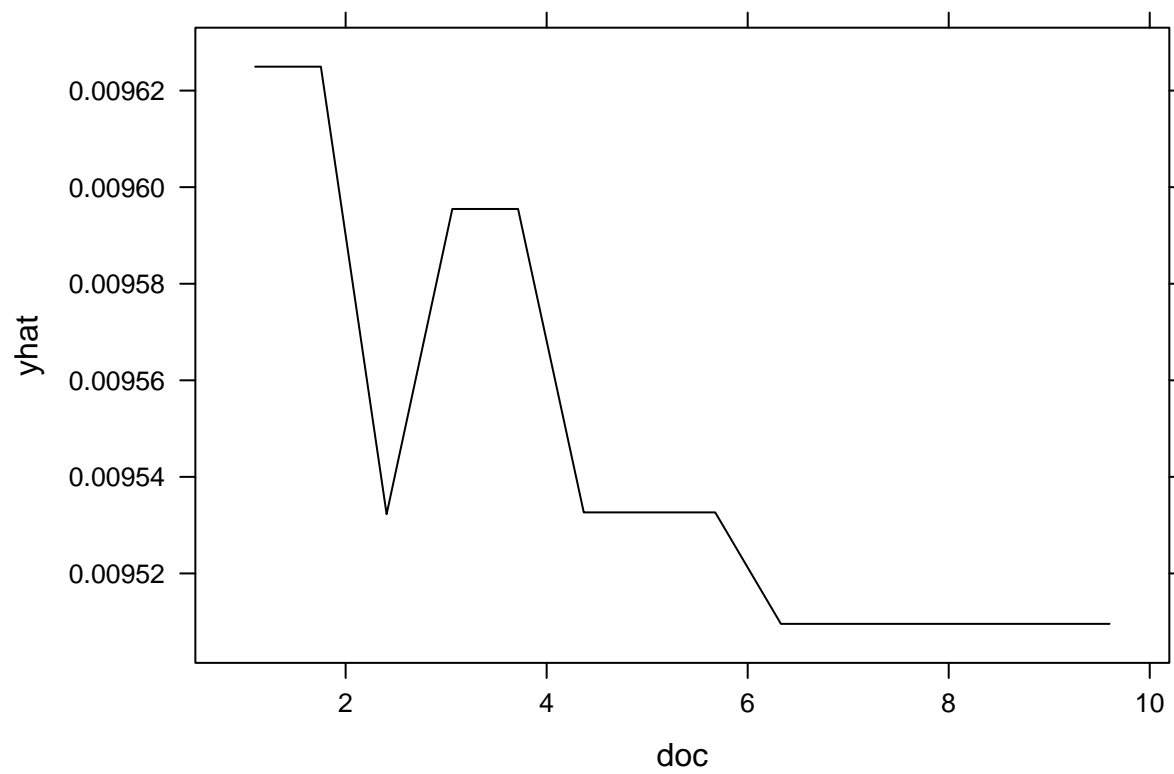
```
partial(cforest.phi.lt, pred.var="prcp.normal", train=modvars.accndvi.philt, type="regression", plot=T)
```



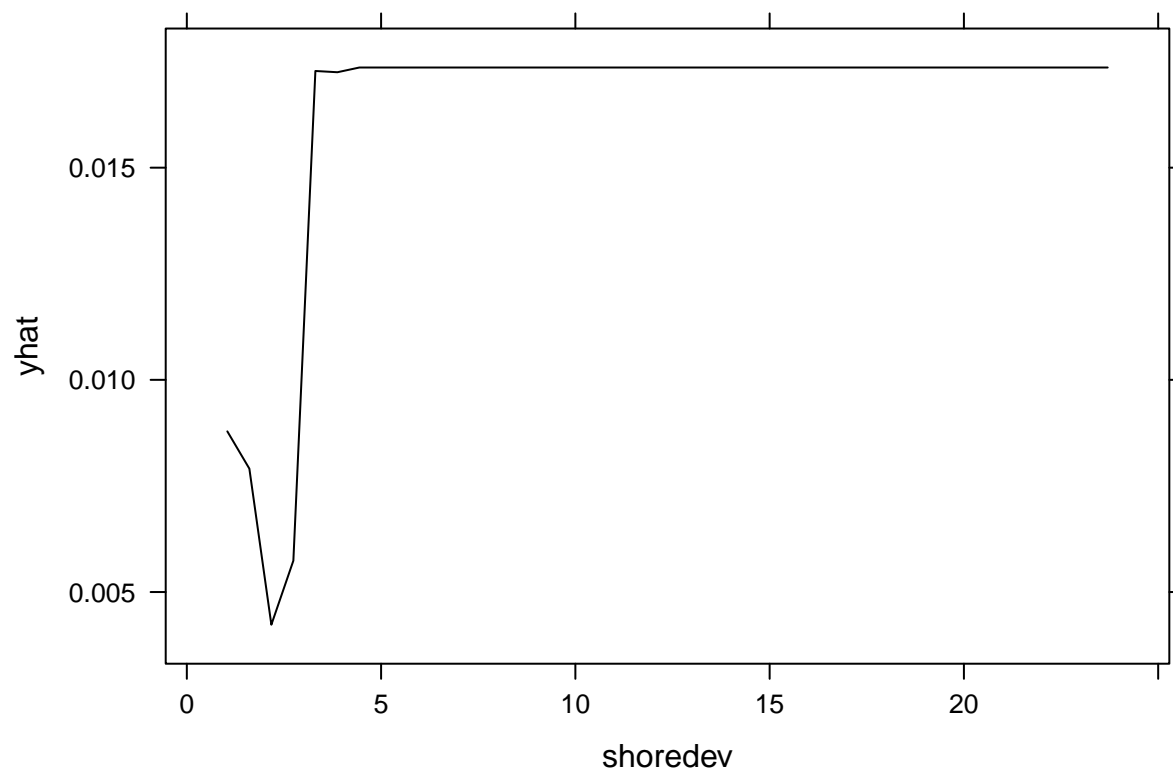
```
partial(cforest.phi.lt, pred.var="pct.wetlands", train=modvars.accndvi.philt, type="regression", plot=T)
```



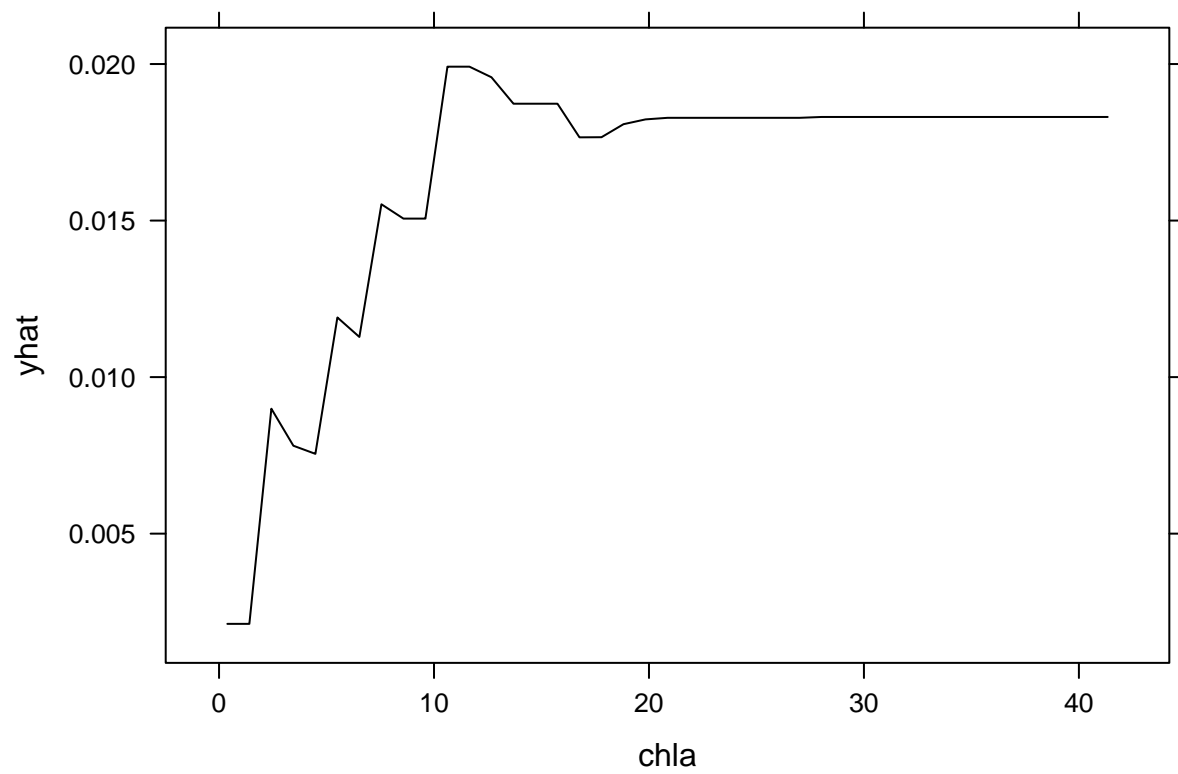
```
partial(cforest.phi.lt, pred.var="doc", train=modvars.accndvi.philt, type="regression", plot=T)
```



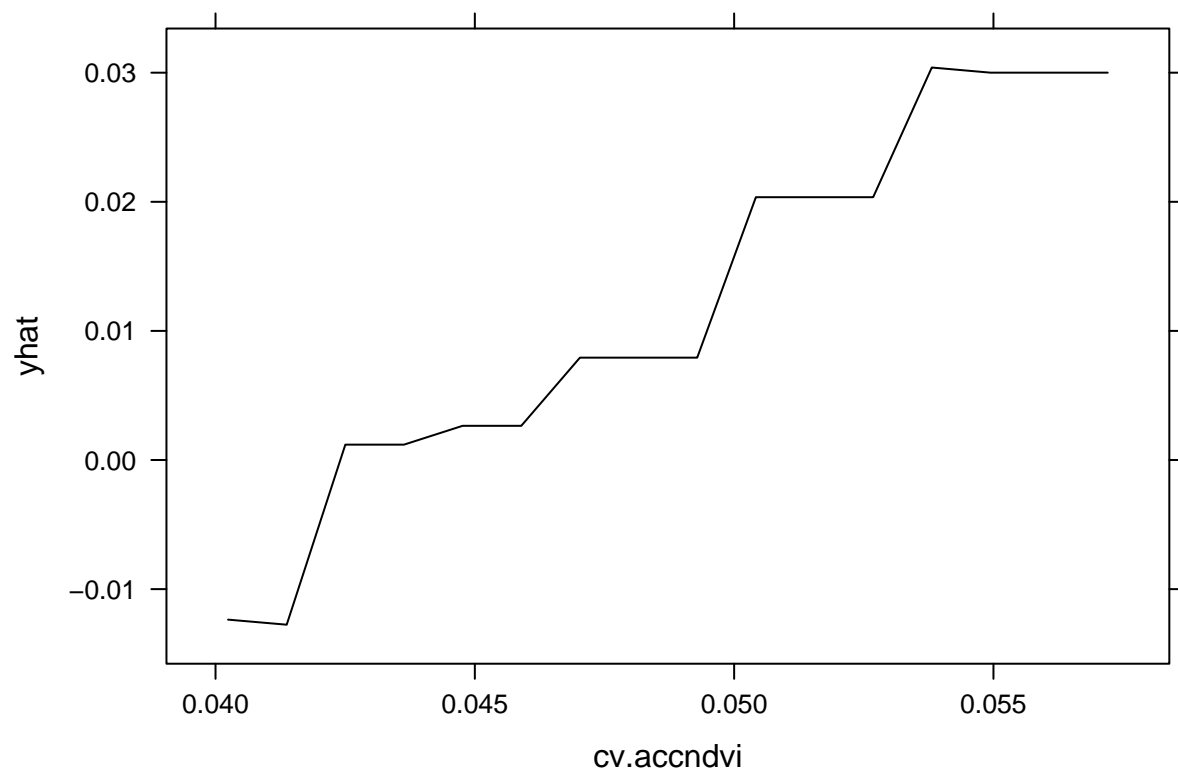
```
partial(cforest.phi.lt, pred.var="shoredv", train=modvars.accndvi.philt, type="regression", plot=T)
```



```
partial(cforest.phi.lt, pred.var="chla", train=modvars.accndvi.philt, type="regression", plot=T)
```

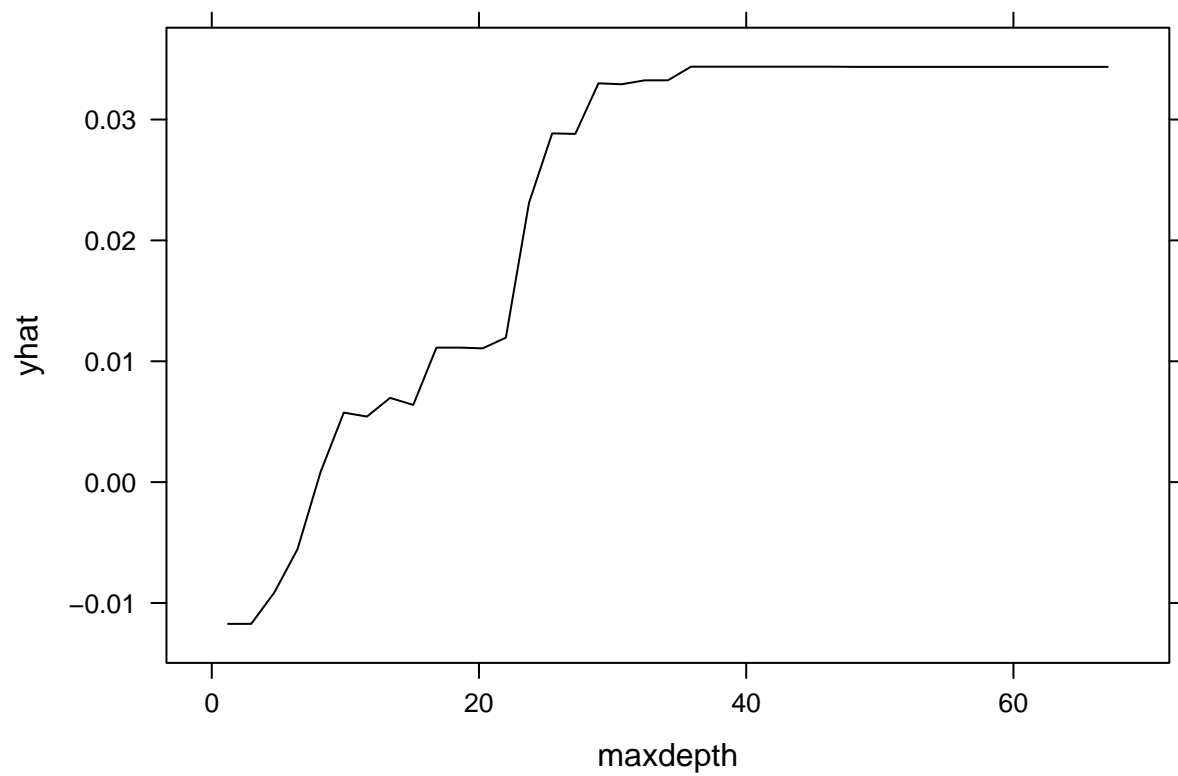


```
partial(cforest.phi.lt, pred.var="cv.accndvi", train=modvars.accndvi.philt, type="regression", plot=T)
```

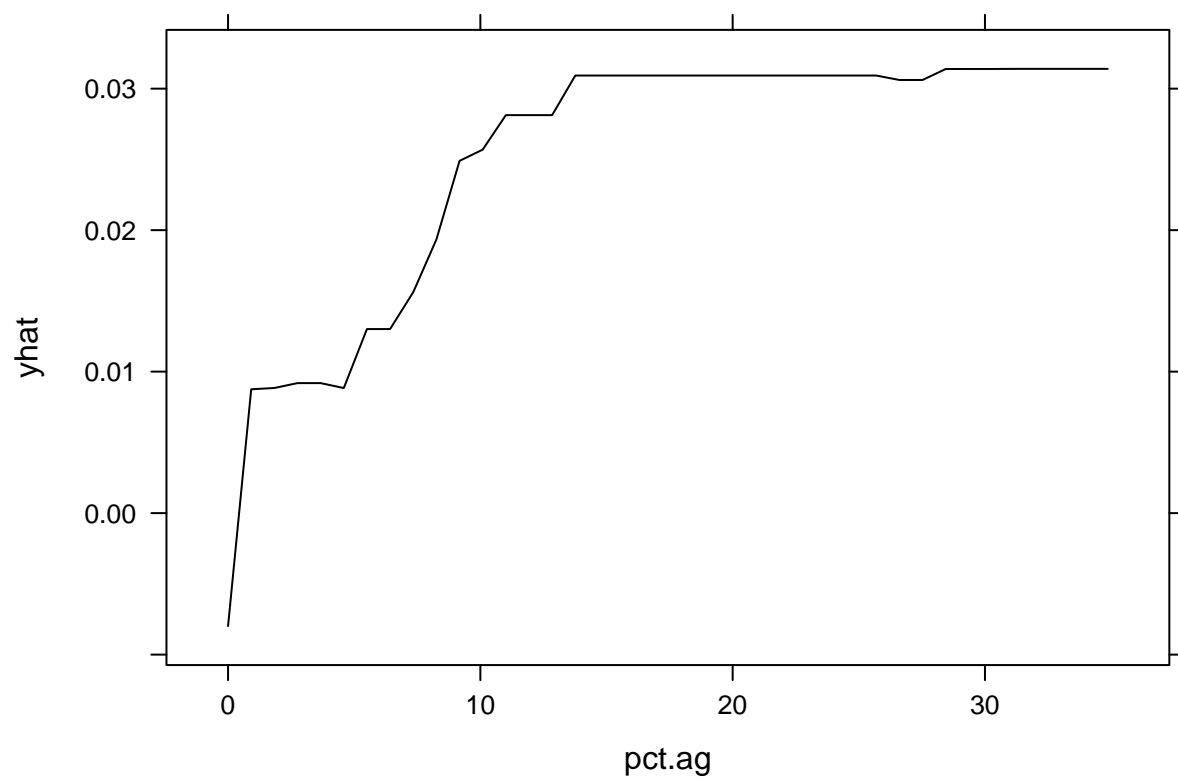


```
partial(cforest.phi.lt, pred.var="maxdepth", train=modvars.accndvi.philt, type="regression", plot=T)
```

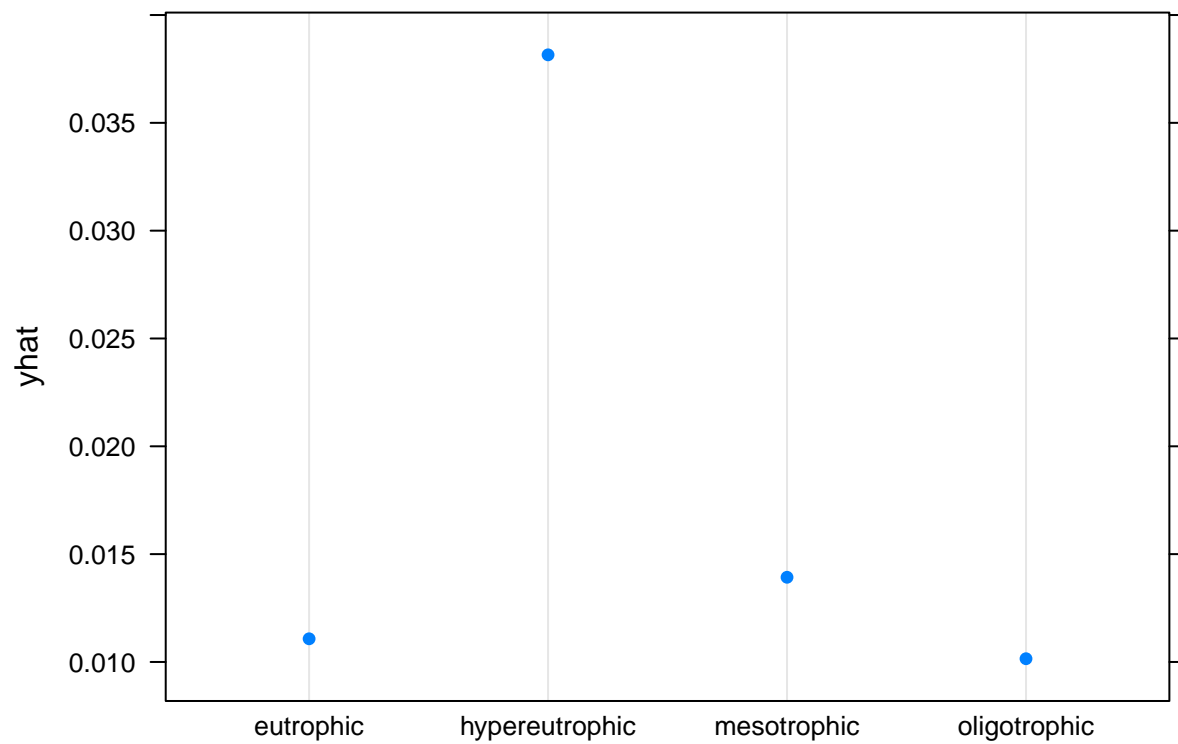




```
partial(cforest.phi.lt, pred.var="pct.ag", train=modvars.accndvi.philt, type="regression", plot=T)
```



```
partial(cforest.phi.lt, pred.var="tsi.cat", train=modvars.accndvi.philt, type="regression", plot=T)
```



```
save.image("~/Box Sync/NSF EAGER Synchrony/Data/RData files/ms1_analyses_complete_20190729.RData")
```