

# 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.RData")

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)

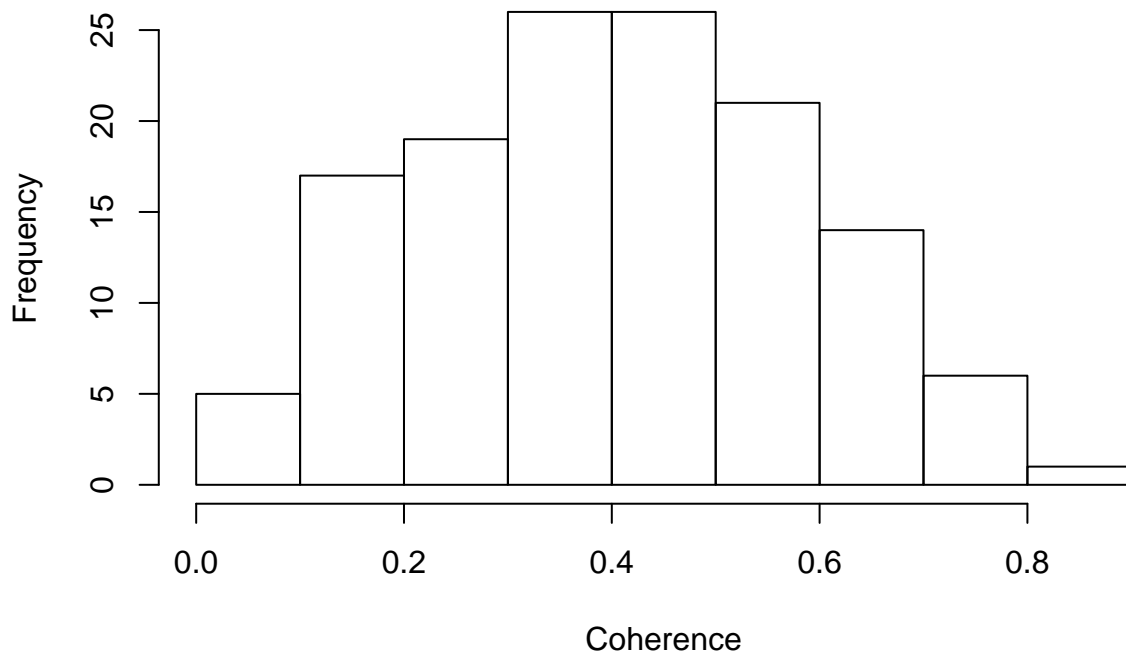
colnames(coh.chlaXaccndvi)<-paste0("accndvi",c("p.ts1","phi.ts1","coh.ts1","p.ts2","phi.ts2","coh.ts2",
colnames(coh.chlaXmaxndvi)<-paste0("maxndvi",c("p.ts1","phi.ts1","coh.ts1","p.ts2","phi.ts2","coh.ts2",

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

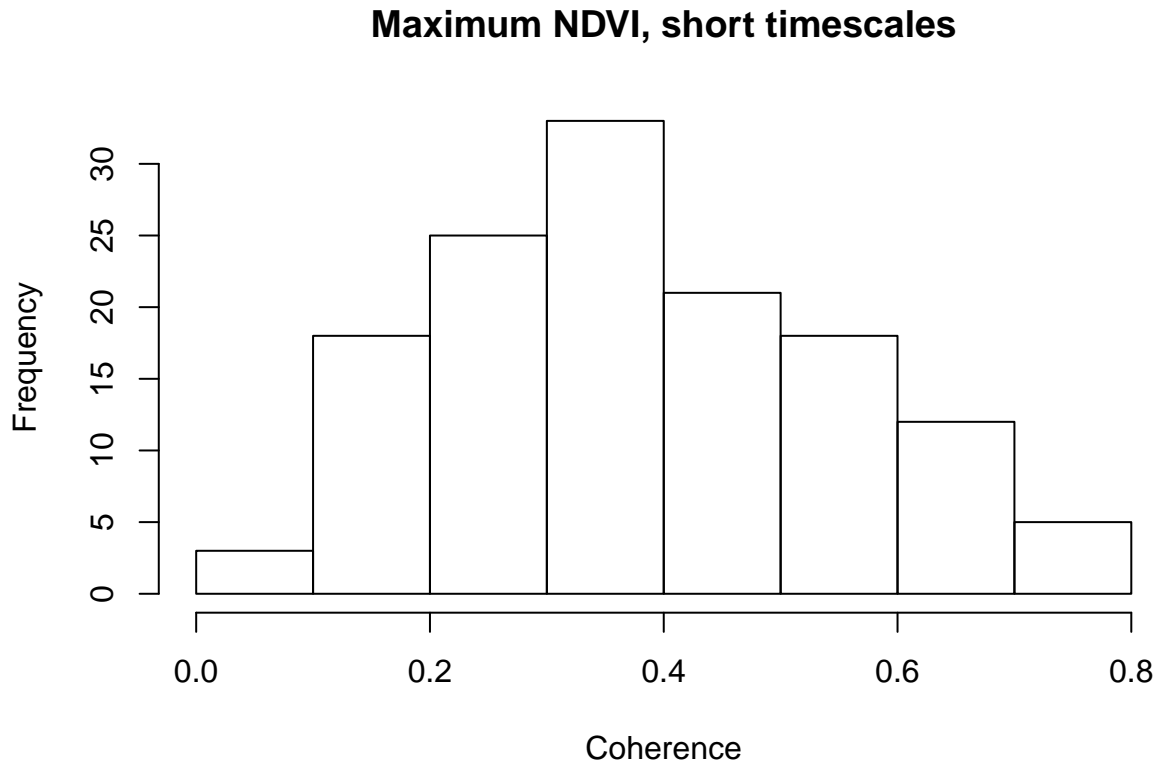
#short timescales
hist(coh.chlaXaccndvi$accndvicoh.ts1, main="Accumulated NDVI, short timescales", xlab="Coherence", ylab="Frequency", col="white", border="black")

```

## 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(coh.chlaXaccndvi$accndviphi.ts1[coh.chlaXaccndvi$accndvip.ts1<alpha]/pi) #only pattern is that la
```

```
## [1] 0.33224850 -0.97156054 -0.04413595 0.56356061 -0.86709075 -0.05260276
## [7] 0.12416199 -0.04172693 0.92429361
```

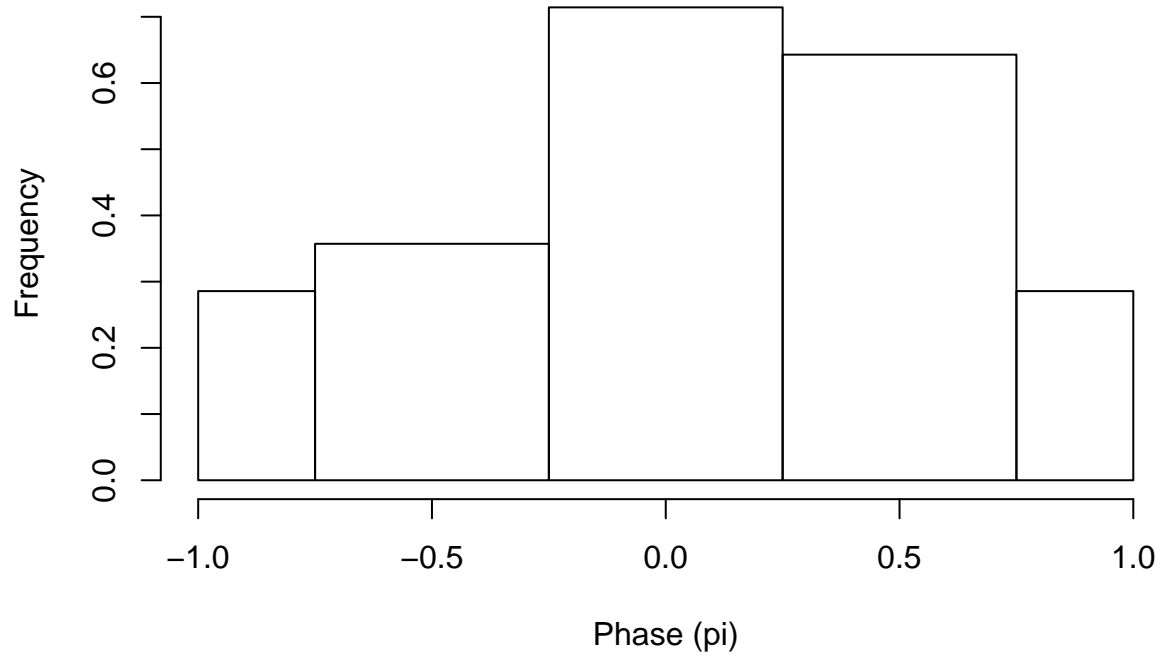
```
print(coh.chlaXmaxndvi$maxndviphi.ts1[coh.chlaXmaxndvi$maxndvip.ts1<alpha]/pi)
```

```
## [1] -0.1573764 -0.8240104 -0.7892870 -0.7185325 -0.9310910 -0.8435071
## [7] -0.2280369 0.5324496
```

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

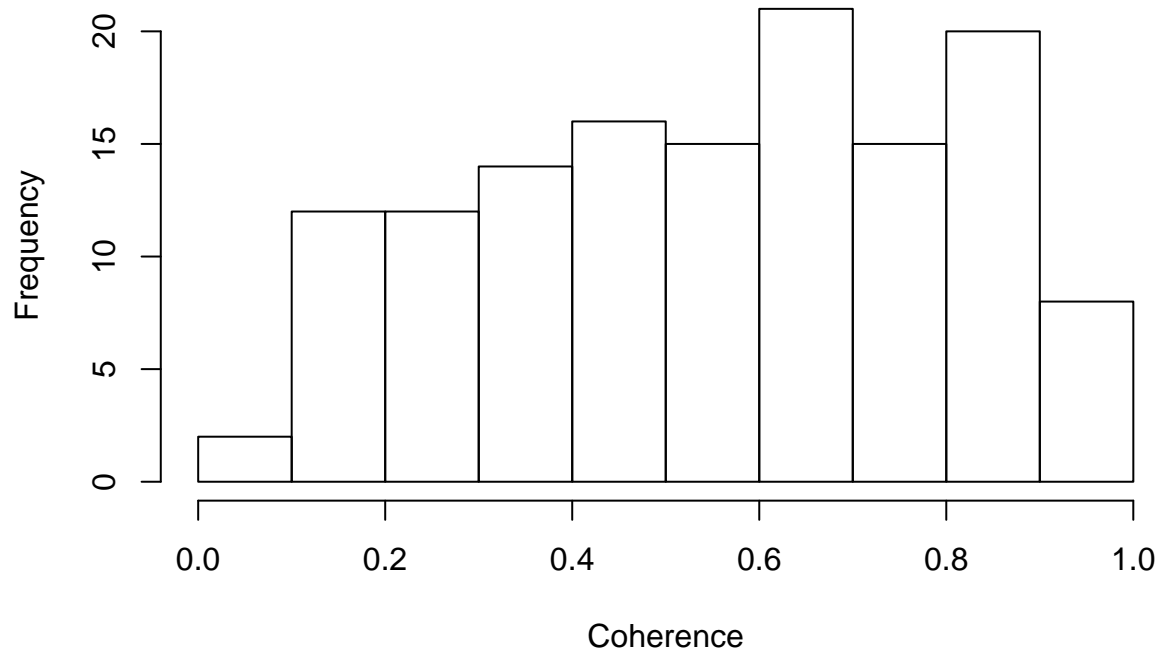
```
hist(coh.chlaXaccndvi$accndviphi.ts1[coh.chlaXaccndvi$accndvip.ts1<0.2]/pi, main="Accumulated NDVI, sho
```

### Accumulated NDVI, short timescales

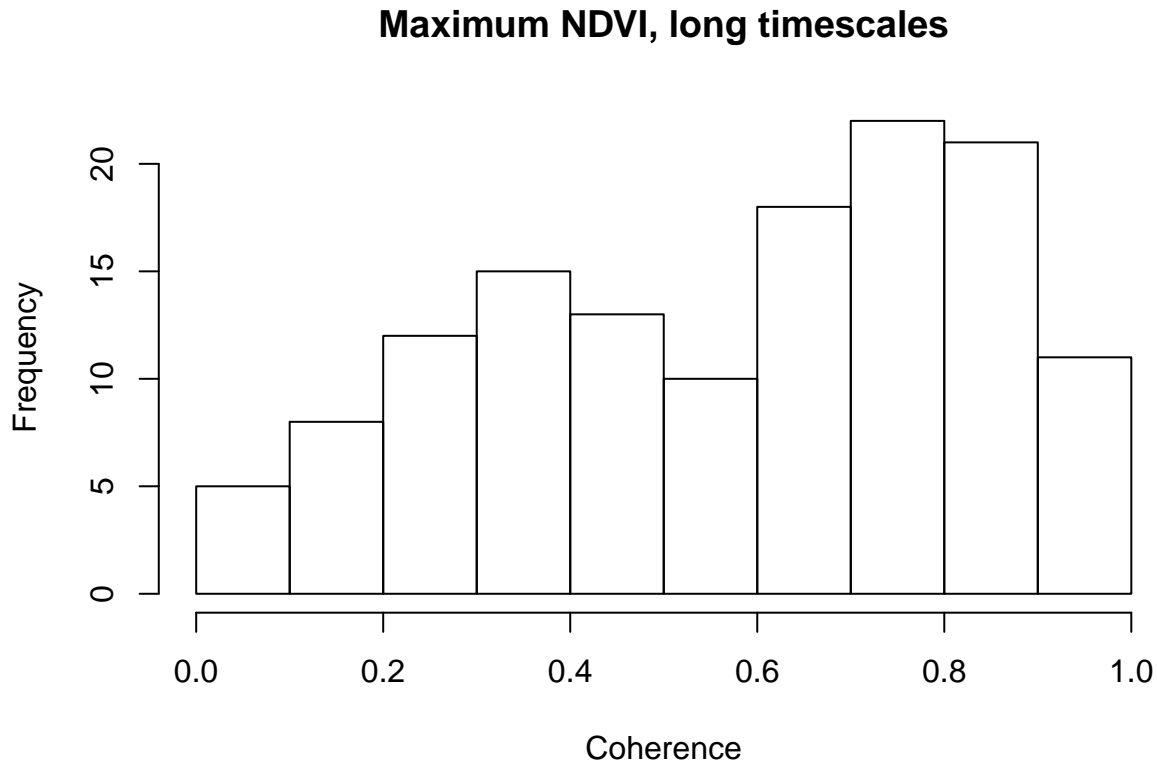


```
#hist(coh.chlaXmaxndvi$maxndviphi.ts1[coh.chlaXmaxndvi$maxndvip.ts1<0.2]/pi, main="Maximum NDVI, short  
#long timescales  
hist(coh.chlaXaccndvi$accndvicoh.ts2, main="Accumulated NDVI, long timescales", xlab="Coherence", ylab=
```

### Accumulated NDVI, long timescales



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



```
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] 0.05925926
```

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

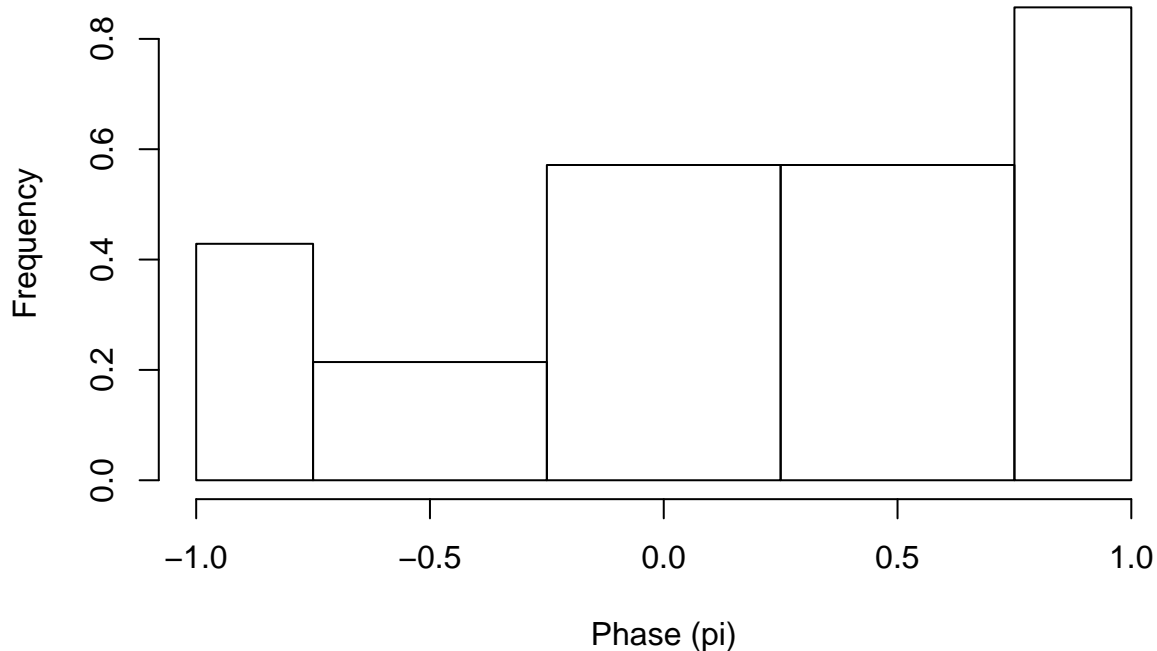
```
## [1] -0.43893809  0.25315167 -0.40935196 -0.04386325 -0.65597599  0.36382168
## [7]  0.89471121
```

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

```
## [1]  0.69982097 -0.97179292 -0.04190360  0.02097044 -0.67004320 -0.58501674
## [7] -0.31373024 -0.33804686
```

```
hist(coh.chlaXaccndvi$accndviphi.ts1[coh.chlaXaccndvi$accndvip.ts2<0.2]/pi, main="Accumulated NDVI, long timescales", xlab="Coherence", ylab="Frequency")
```

## Accumulated NDVI, long timescales



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

```
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
```

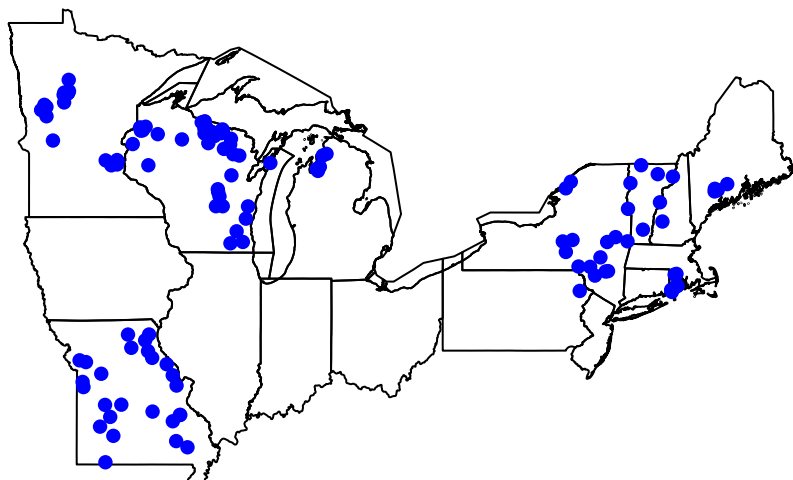
```
getstates<-c("Minnesota", "Iowa", "Wisconsin", "Illinois", "Missouri", "Michigan", "Indiana", "Ohio", "Louisiana", "Alabama", "Georgia", "Florida", "South Carolina", "North Carolina", "Virginia", "Maryland", "Delaware", "Pennsylvania", "New York", "New Jersey", "Connecticut", "Massachusetts", "Rhode Island", "Vermont", "New Hampshire", "Maine", "Hawaii")
```

```
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



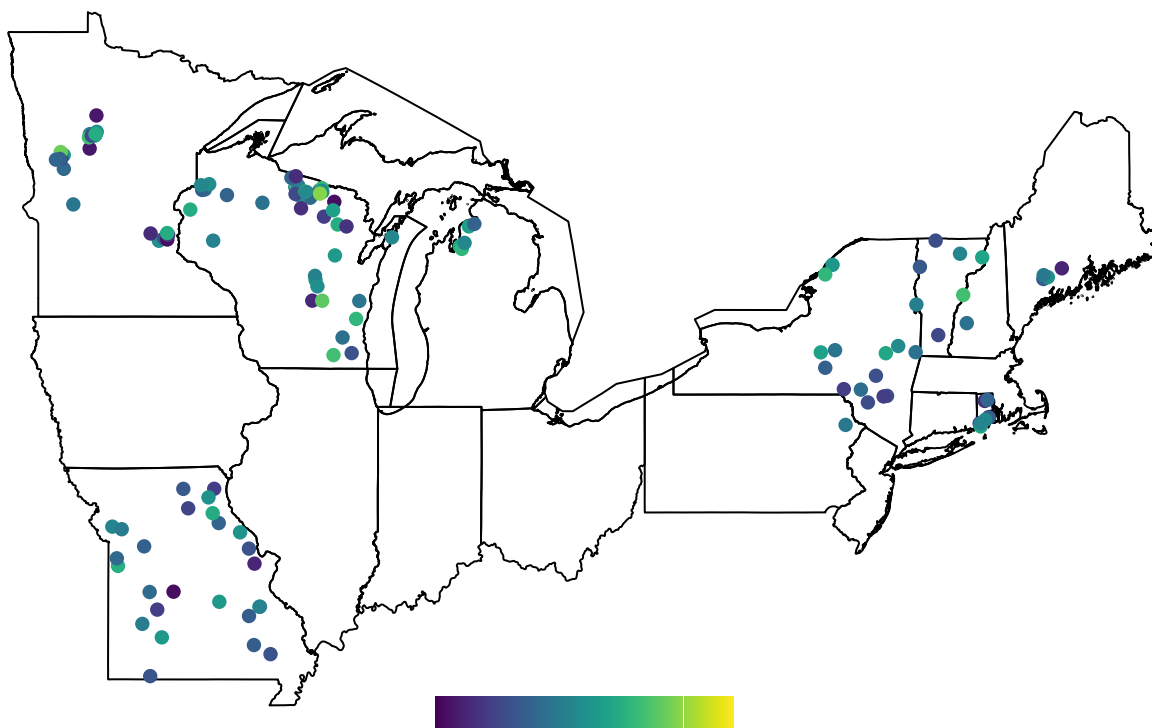
```
cohplotdata<-left_join(analysislakes$lakeinfo, coh.chlaXaccndvi, by="lagoslakeid")

pal<-viridis(100)

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

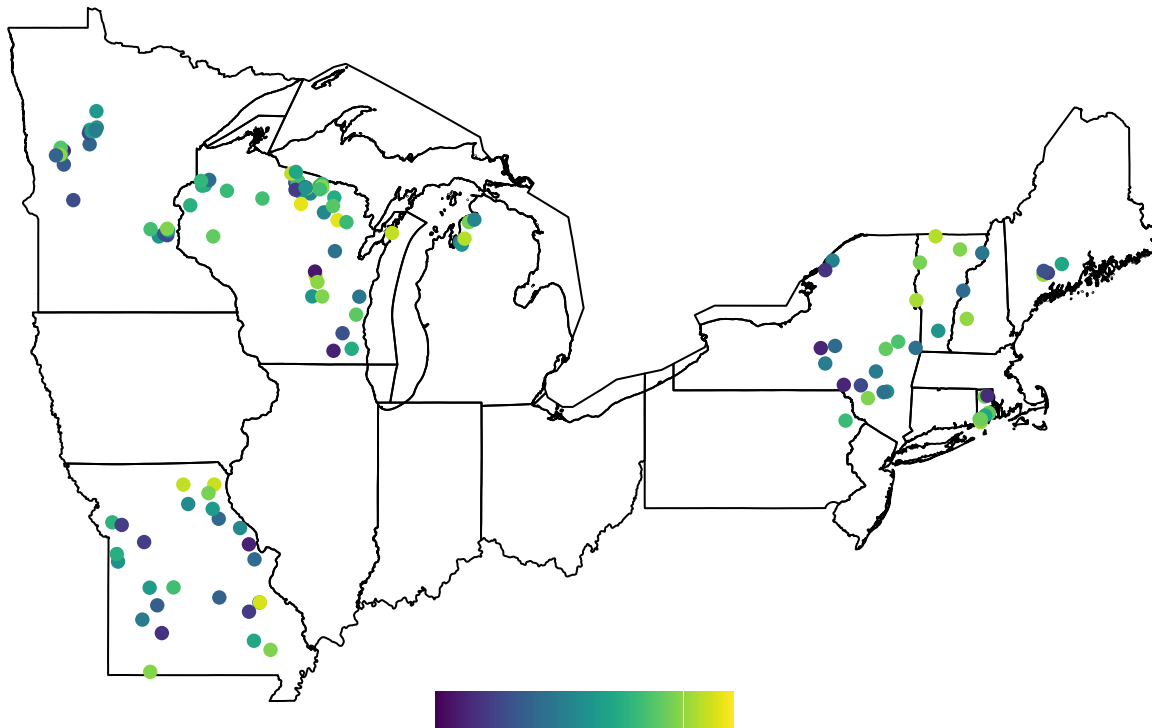
plot(lagosstates, main="Lakes by short timescale coherence")
points(cohplotdata$nhd_long, cohplotdata$nhd_lat, pch=16, cex=1, col=pal[round(cohplotdata$accndvicoh.t.
colorbar.plot(x=mean(par("usr")[1:2]),y=par("usr")[3],strip=1:100,col=pal,horizontal = T)
```

## 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_pc
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]))

#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 mo

#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)

#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)

#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"))

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, 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.accndvi, 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)

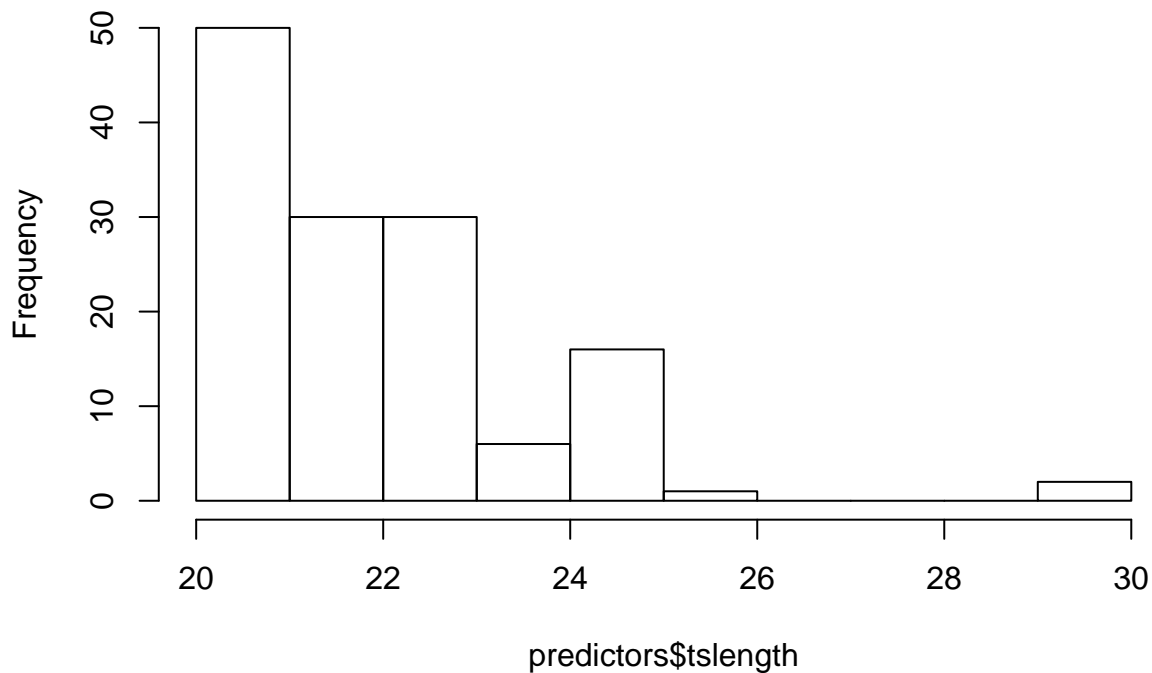
## 'data.frame':   135 obs. of  22 variables:
## $ lagoslakeid      : num  211 249 618 906 969 ...
## $ gnis_name        : chr   NA NA "Butternut Lake" "Sparkling Lake" ...
## $ nhd_lat          : num  44.5 43.7 45.9 46 45.8 ...
## $ nhd_long         : num  -73.3 -73.4 -89 -89.7 -89.3 ...
## $ lake_area_ha     : 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 ...
## $ nhd_fcode        : int  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 ...
## $ 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.accndvi      : num  0.0572 0.0542 0.0443 0.0561 0.0417 ...
```

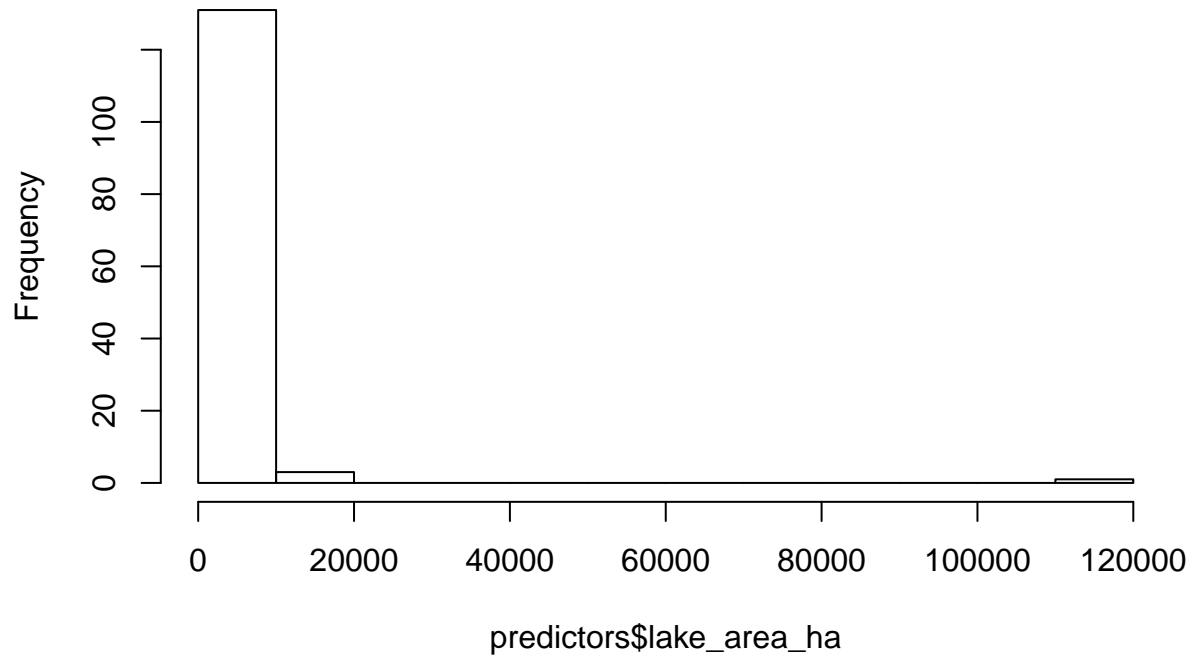
```
hist(predictors$tslength)
```

**Histogram of predictors\$tslength**



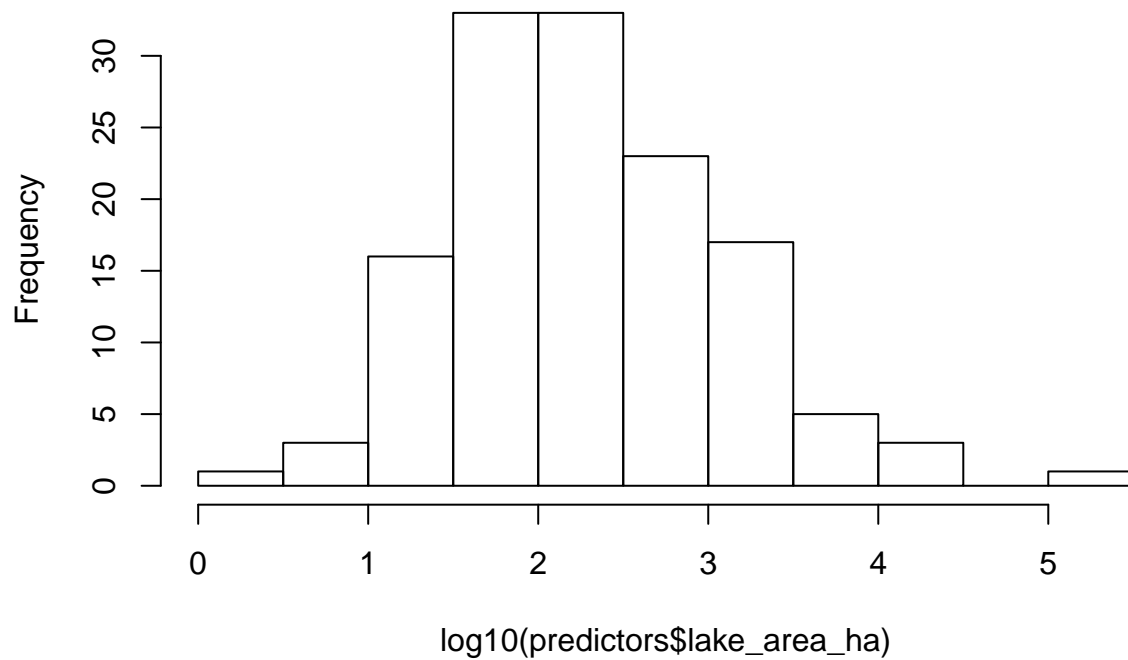
```
hist(predictors$lake_area_ha)
```

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



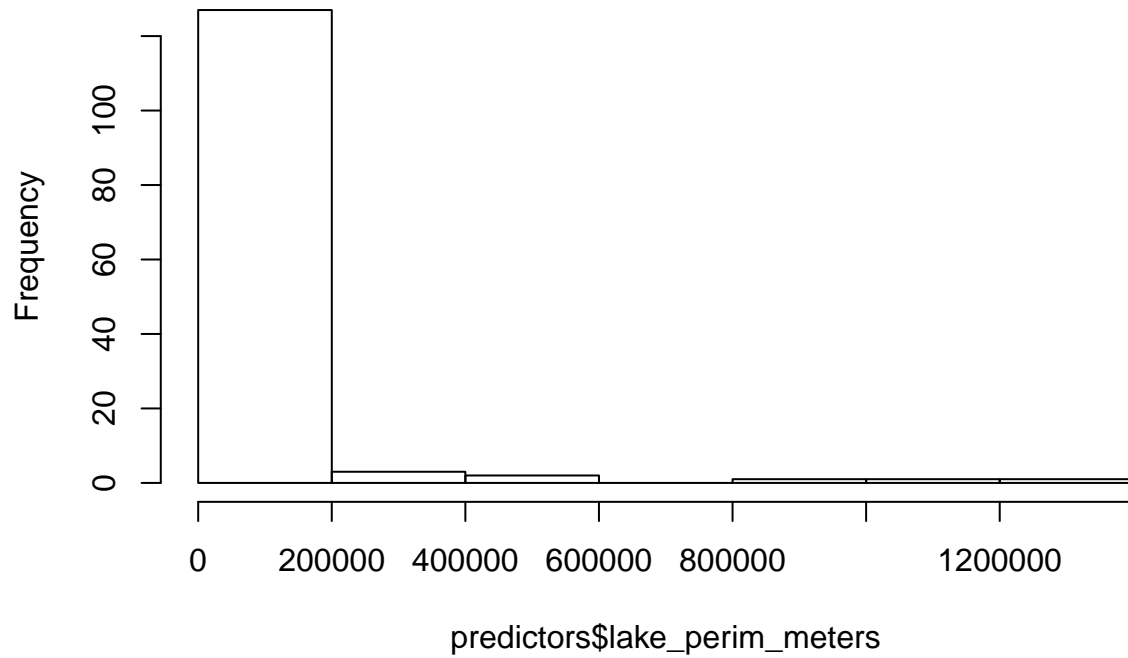
```
hist(log10(predictors$lake_area_ha))
```

**Histogram of log10(predictors\$lake\_area\_ha)**



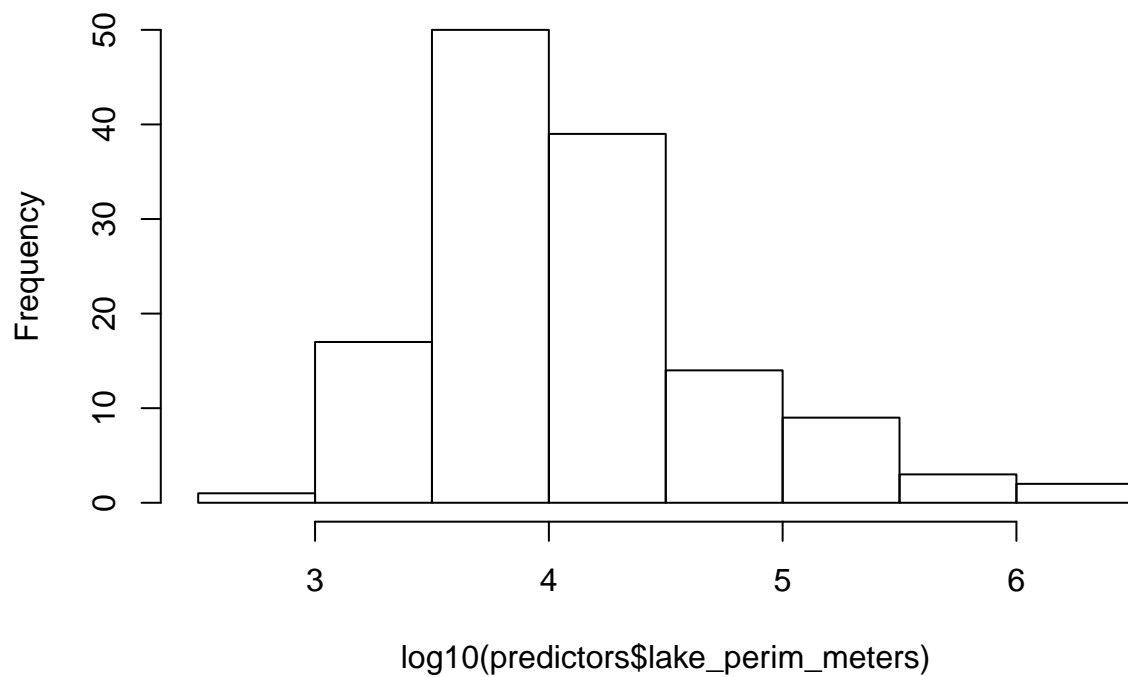
```
hist(predictors$lake_perim_meters)
```

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



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

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



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

```
##
```

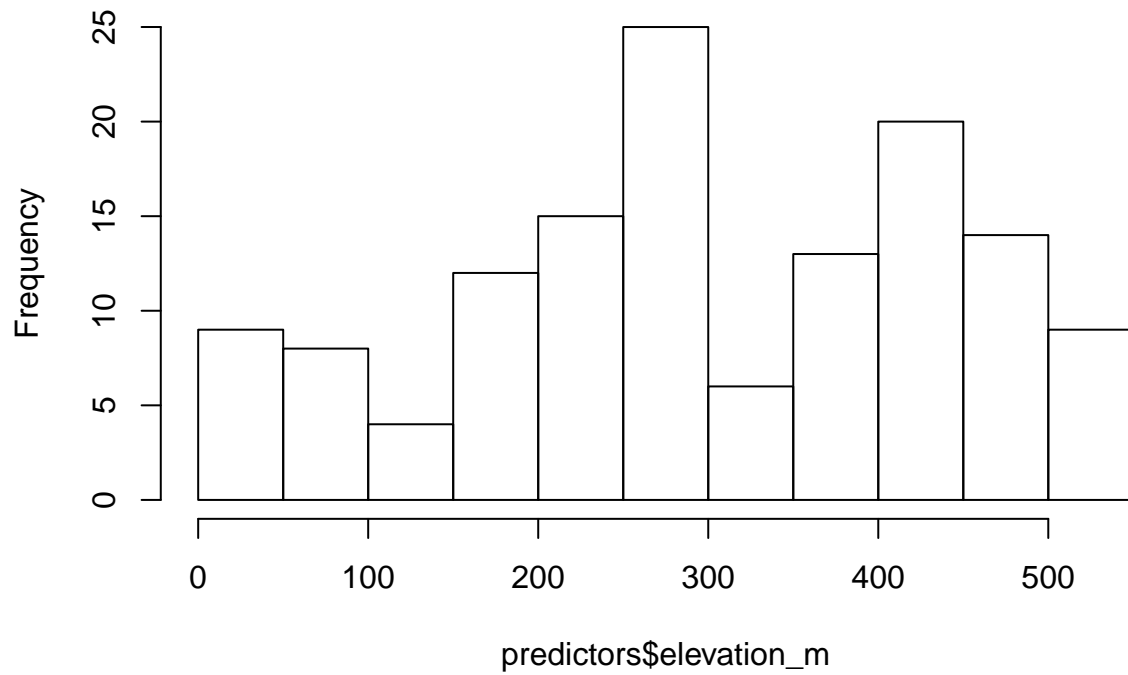
```
## 39000 39004 39009 39010 39012 43601
##      1   110    14     3     6     1
```

```
table(predictors$hul2_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
```

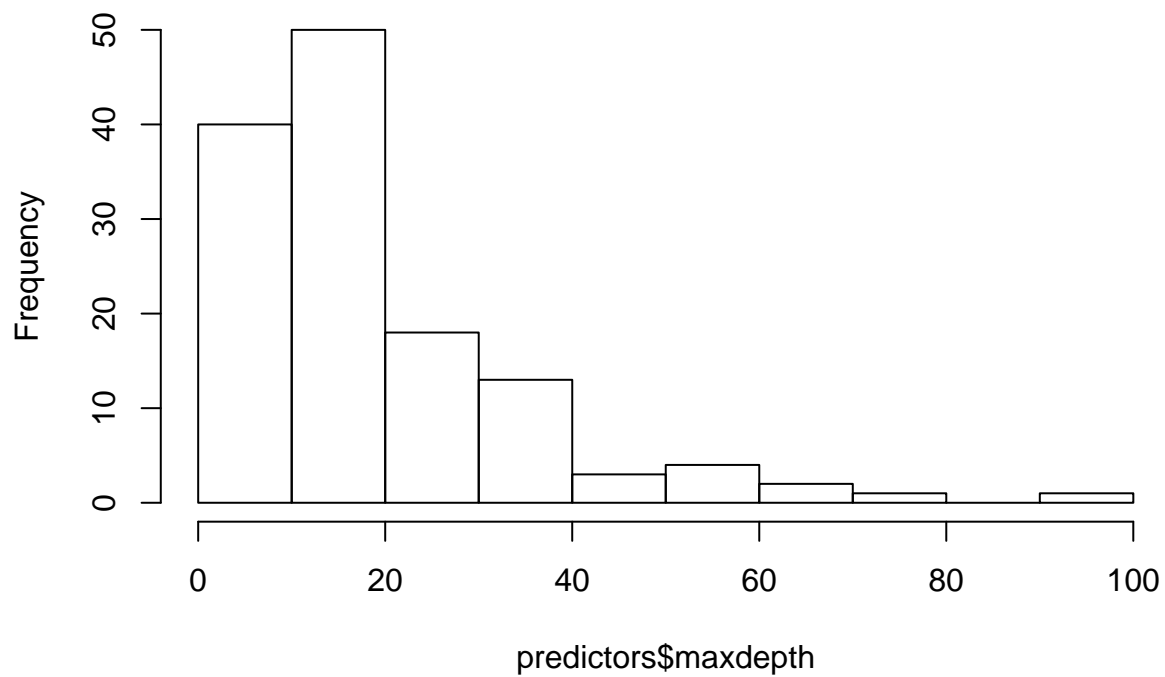
```
hist(predictors$elevation_m)
```

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



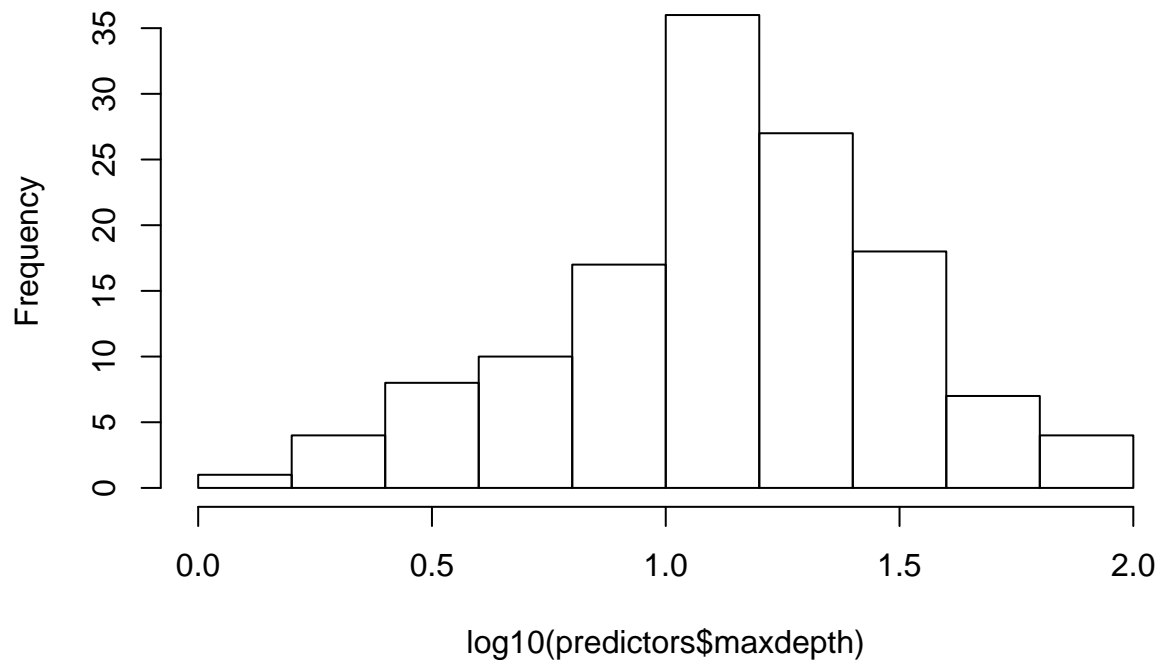
```
hist(predictors$maxdepth)
```

**Histogram of predictors\$maxdepth**



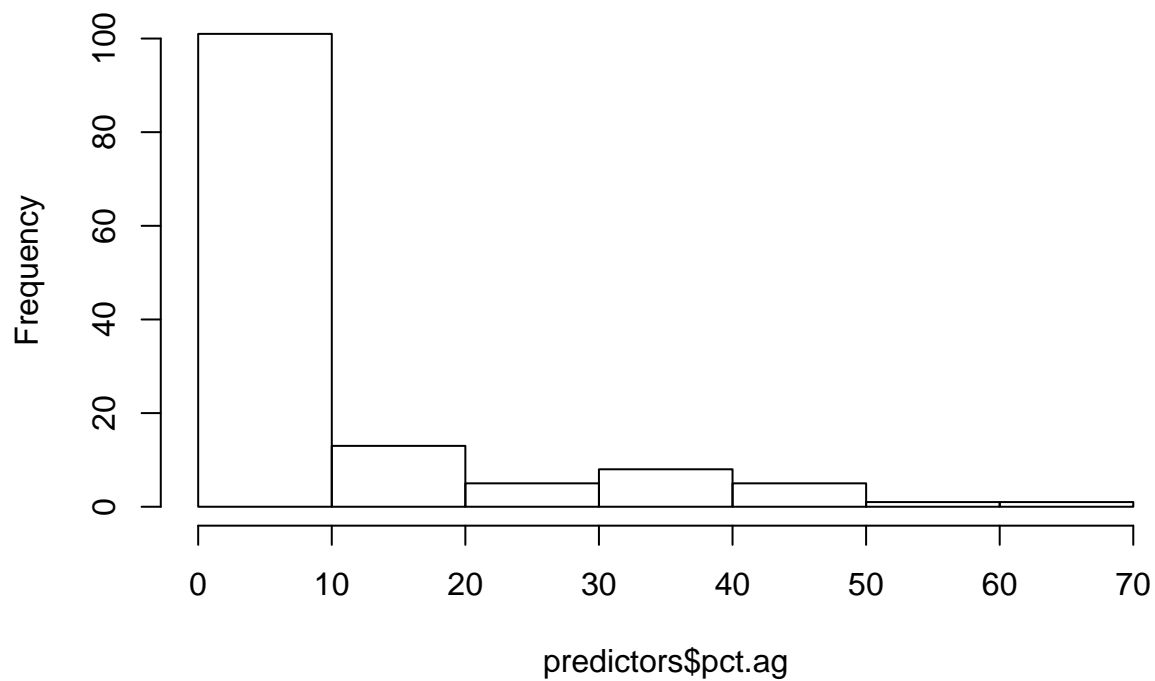
```
hist(log10(predictors$maxdepth))
```

**Histogram of  $\log_{10}(\text{predictors}\$maxdepth)$**



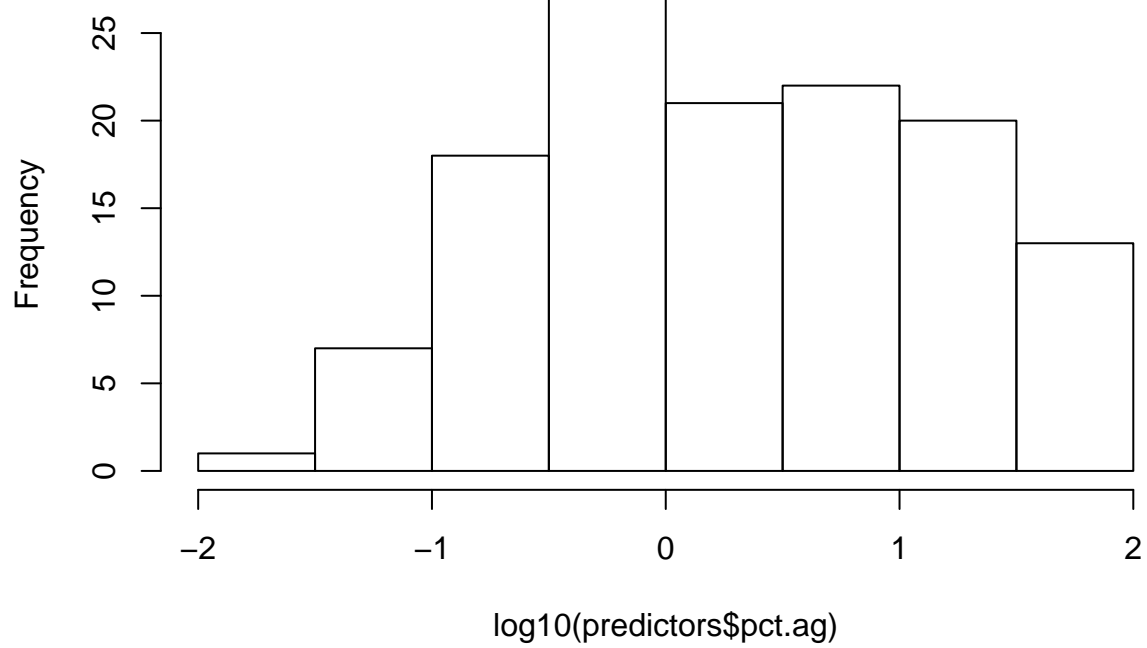
```
hist(predictors$pct.ag)
```

**Histogram of  $\text{predictors}\$pct.ag$**



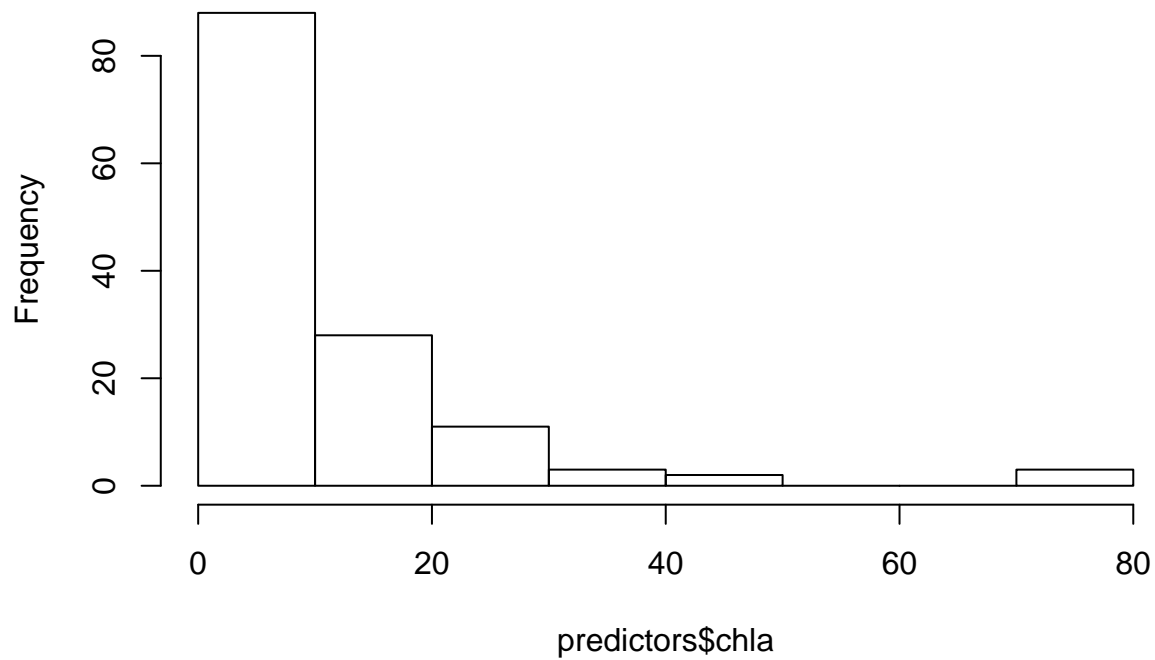
```
hist(log10(predictors$pct.ag))
```

**Histogram of  $\log_{10}(\text{predictors}\$pct.ag)$**



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

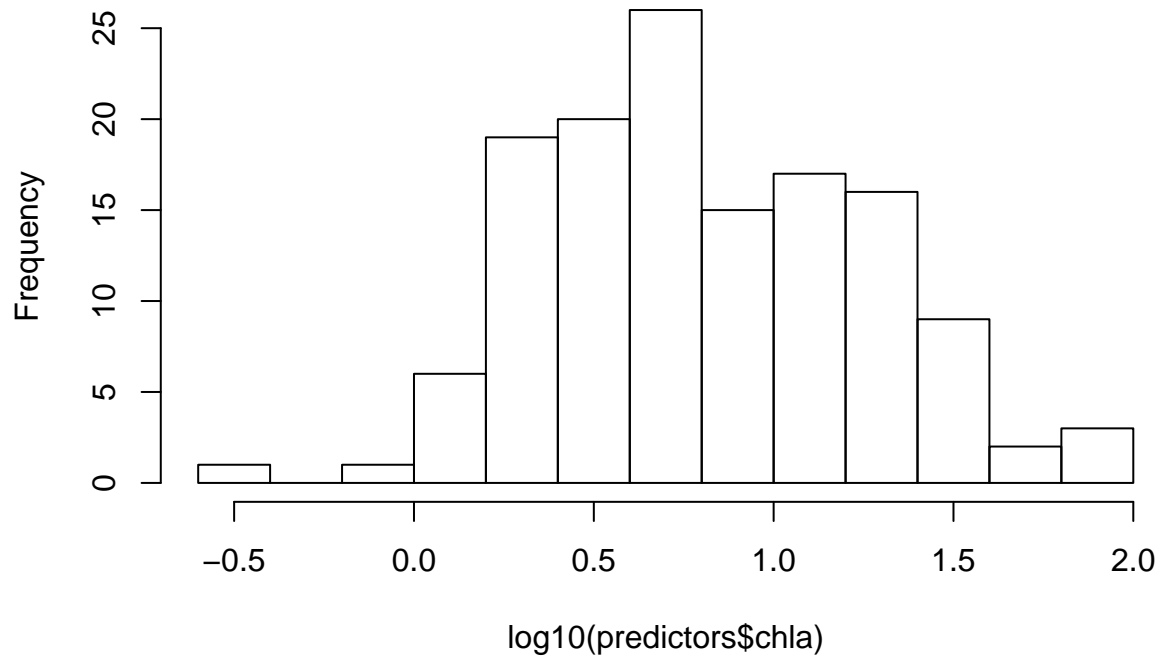
**Histogram of predictors\$chla**



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

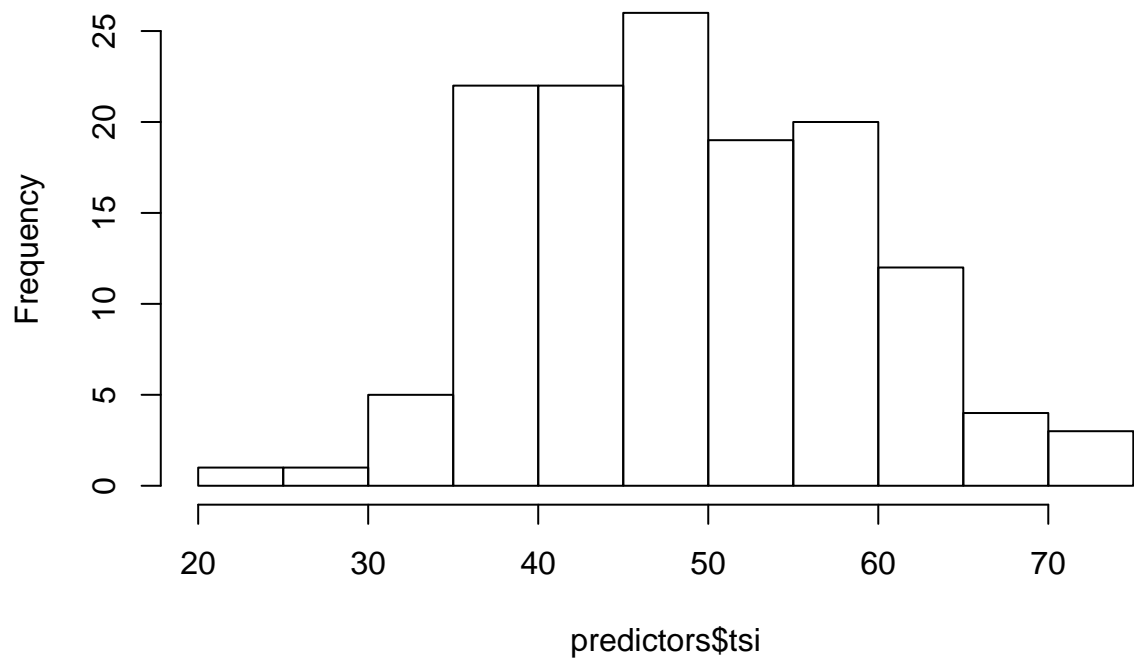


**Histogram of  $\log_{10}(\text{predictors}\$chla)$**



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

**Histogram of  $\text{predictors}\$tsi$**

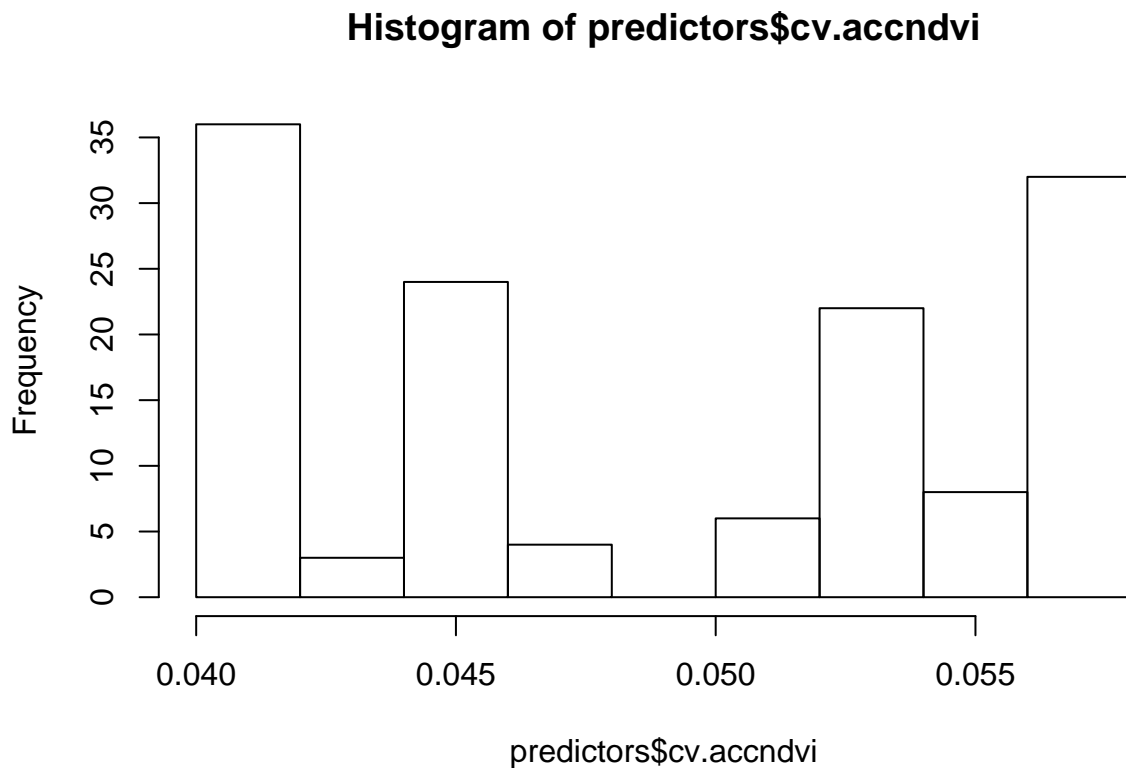


```
table(predictors$tsi.cat)
```

```
##
```

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

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



```
predictors$log10_lake_area_ha<-log10(predictors$lake_area_ha)
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,]
```

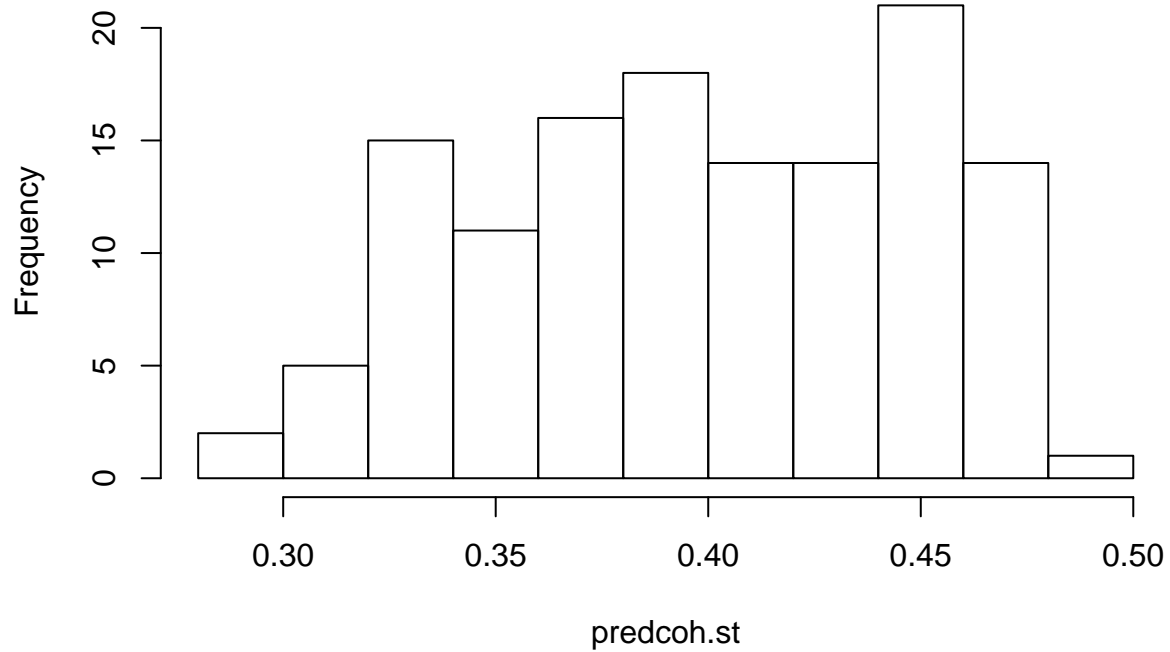
```
library(partykit)
```

```
## Loading required package: libcoin
```

```
## Loading required package: mvtnorm
```

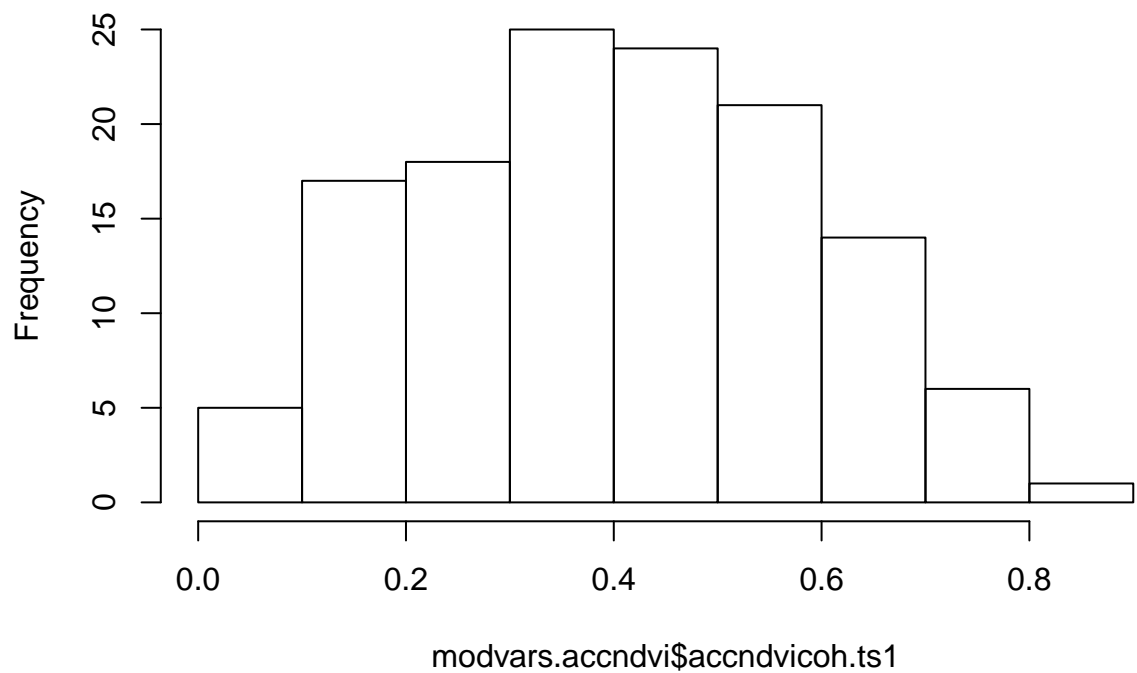
```
cforest.st<-cforest(accndvicoh.ts1 ~ lake_area_ha + lake_perim_meters + maxdepth + pct.ag + chla + tsi +
predcoh.st<-predict.cforest(cforest.st, newdata=modvars.accndvi)
hist(predcoh.st)
```

**Histogram of predcoh.st**



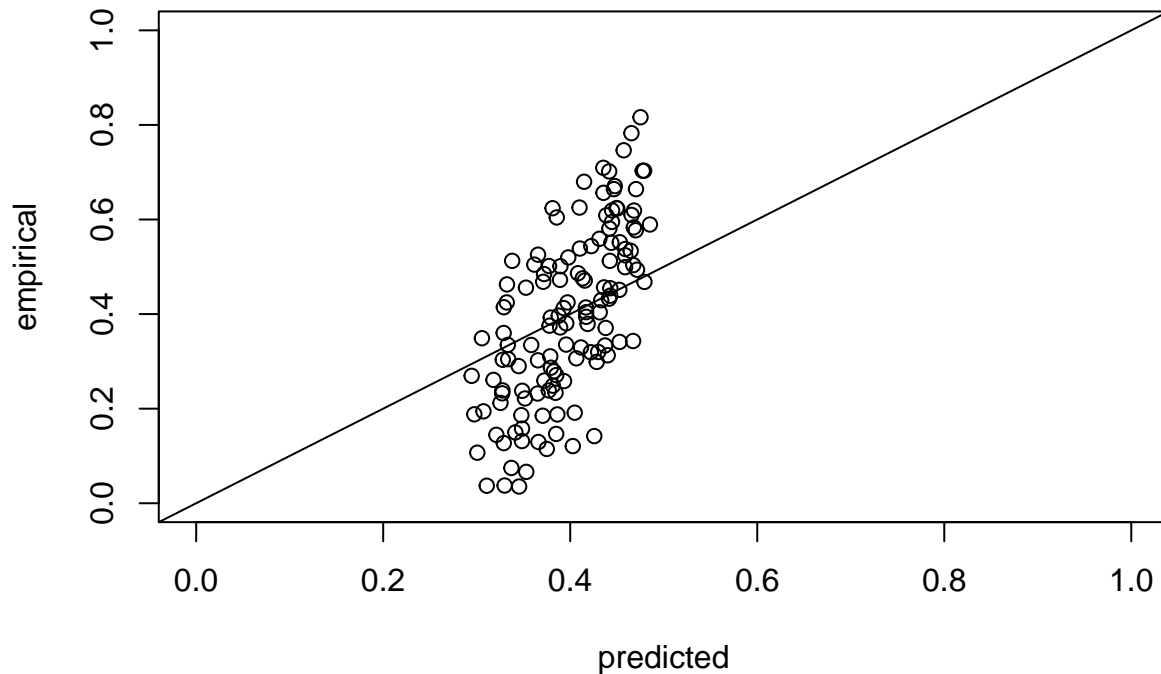
```
hist(modvars.accndvi$accndvicoh.ts1)
```

**Histogram of modvars.accndvi\$accndvicoh.ts1**



```
plot(predcoh.st, modvars.accndvi$accndvicoh.ts1, xlab="predicted", ylab="empirical", main="Coherence, s  
      xlim=c(0,1), ylim=c(0,1))  
abline(a=0,b=1)
```

## Coherence, short ts



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

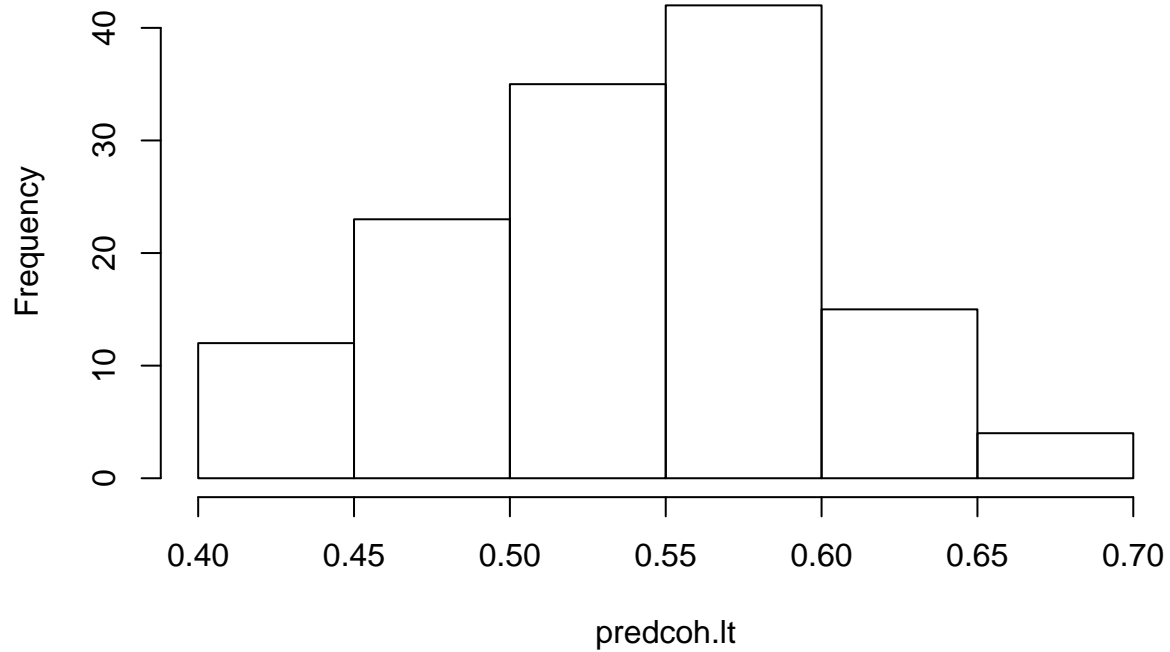
```
##
## Pearson's product-moment correlation
##
## data: predcoh.st and modvars.accndvi$accndvicoh.ts1
## t = 10.618, df = 129, p-value < 2.2e-16
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.5792340 0.7648444
## sample estimates:
##      cor
## 0.6829101
```

```
varimp.coh.st<-varimp.cforest(cforest.st)
print(varimp.coh.st)
```

```
##      lake_area_ha lake_perim_meters      maxdepth      pct.ag
## 1.575424e-03    8.503891e-04    8.139477e-04    1.220314e-03
##          chla          tsi      hu4_zoneid      cv.accndvi
## 5.259568e-04    3.057602e-04   -6.340181e-05   -6.894471e-04
```

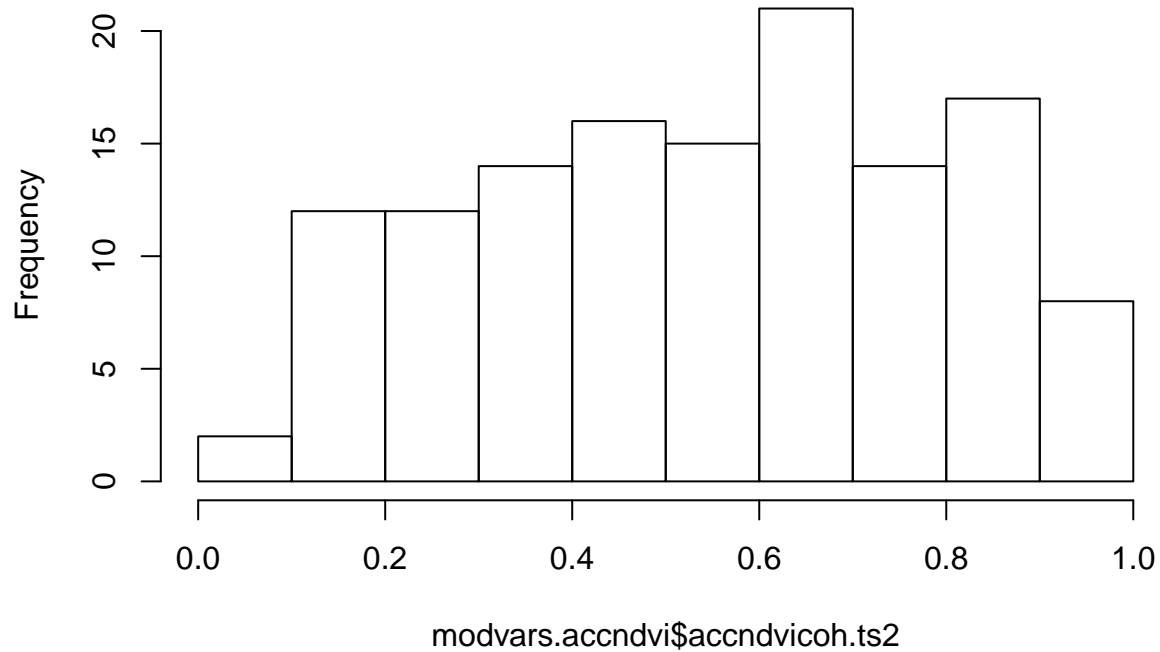
```
cforest.lt<-cforest(accndvicoh.ts2 ~ lake_area_ha + lake_perim_meters + maxdepth + pct.ag + chla + tsi + hu4_zoneid + cv.accndvi)
predcoh.lt<-predict.cforest(cforest.lt, newdata=modvars.accndvi)
hist(predcoh.lt)
```

**Histogram of predcoh.lt**



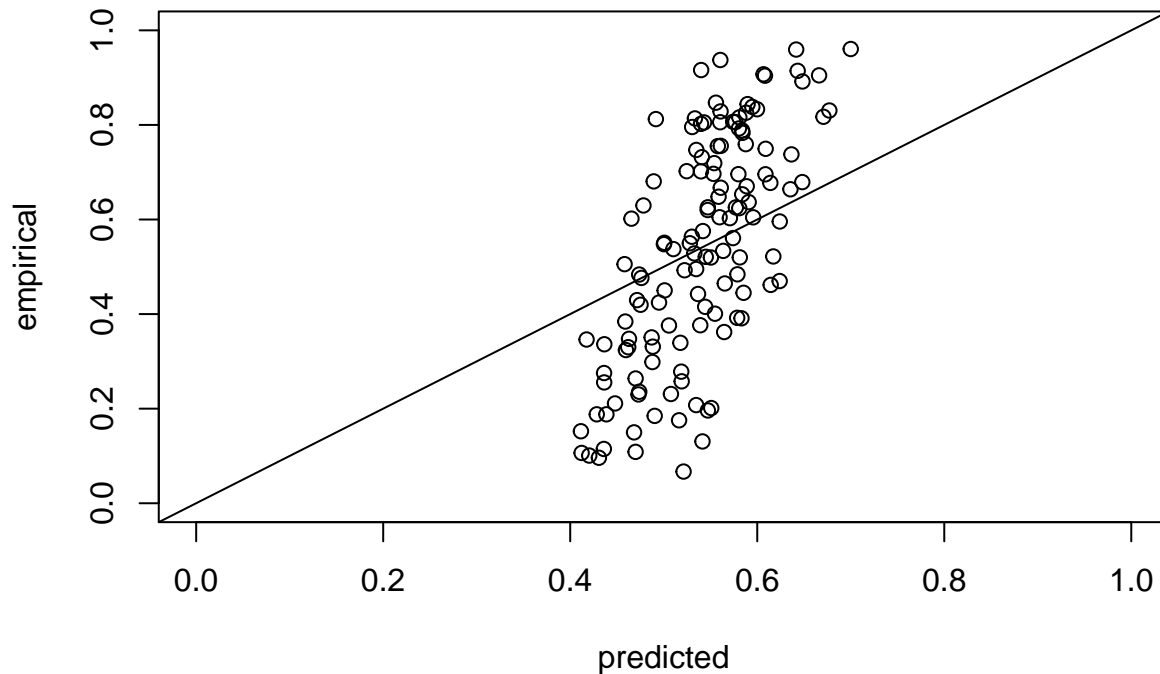
```
hist(modvars.accndvi$accndvicoh.ts2)
```

**Histogram of 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)
```

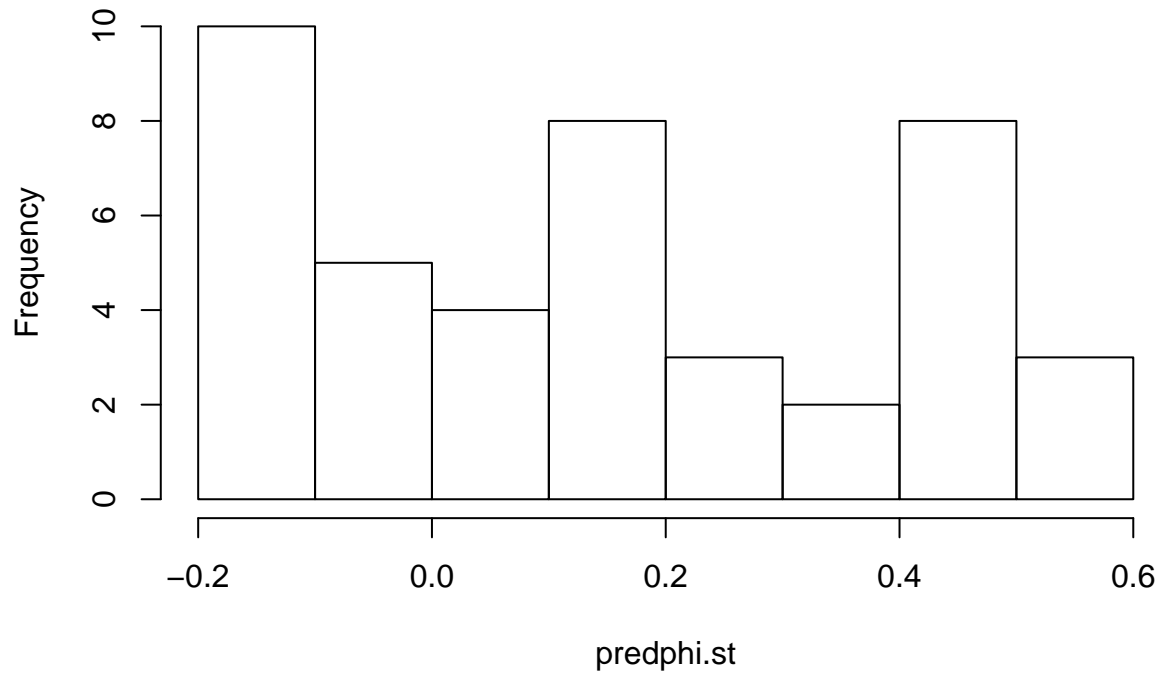
```
##
## Pearson's product-moment correlation
##
## data: predcoh.lt and modvars.accndvi$accndvicoh.ts2
## t = 11.025, df = 129, p-value < 2.2e-16
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.5962244 0.7754058
## sample estimates:
##      cor
## 0.6965185
```

```
varimp.coh.lt<-varimp.cforest(cforest.lt)
print(varimp.coh.lt)
```

```
##      lake_area_ha lake_perim_meters      maxdepth      pct.ag
## -0.0011828020    -0.0003966179    -0.0014873815    0.0002196945
##           chla           tsi      hu4_zoneid      cv.accndvi
##  0.0007098654    -0.0002411120    -0.0005973412    0.0004066143
```

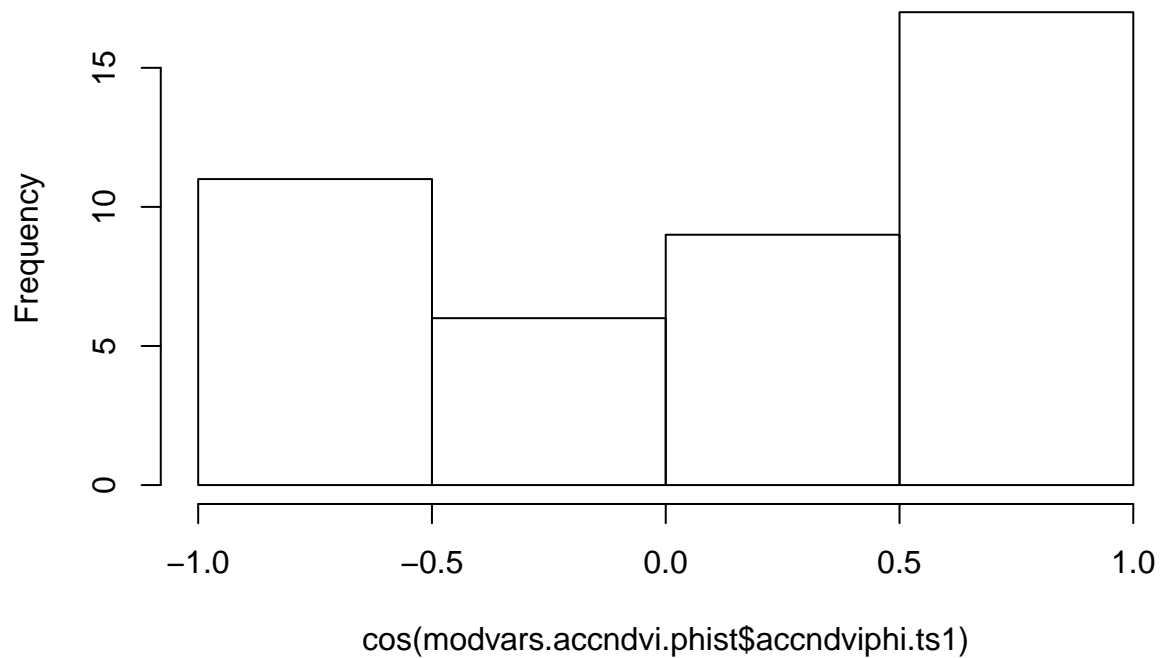
```
cforest.phi.st<-cforest(cos(accndviphi.ts1) ~ lake_area_ha + lake_perim_meters + maxdepth + pct.ag + ch
                        data=modvars.accndvi.phist)
predphi.st<-predict.cforest(cforest.phi.st, newdata=modvars.accndvi.phist)
hist(predphi.st)
```

### Histogram of predphi.st



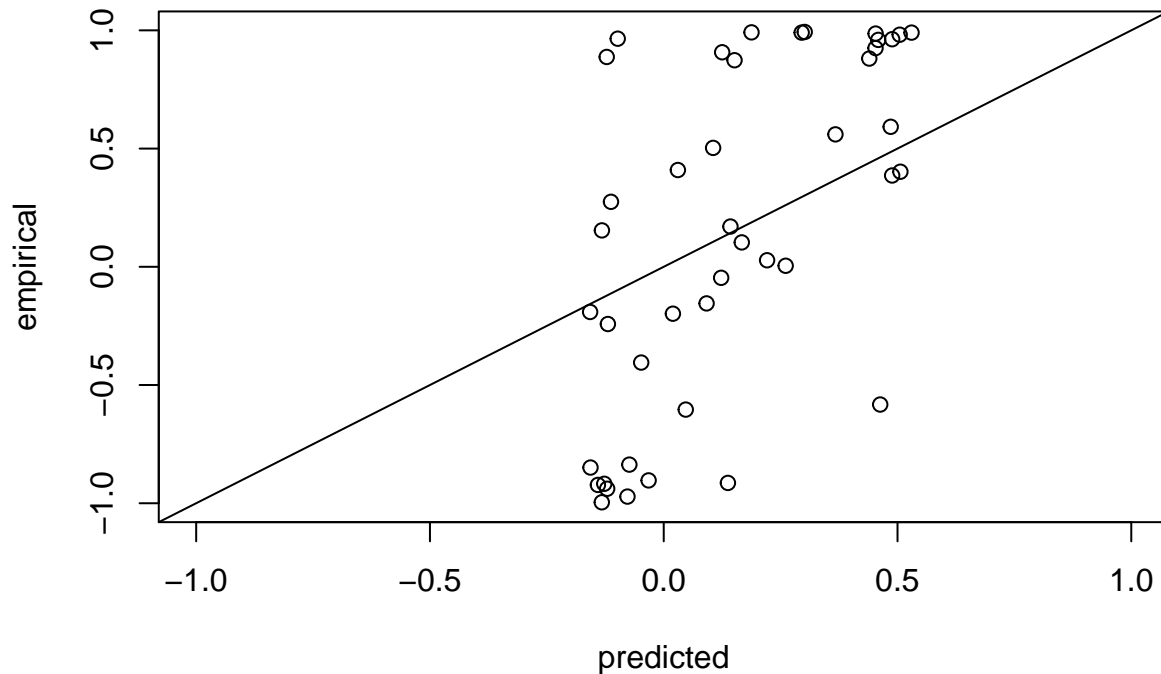
```
hist(cos(modvars.accndvi.phist$accndviphi.ts1))
```

### Histogram of cos(modvars.accndvi.phist\$accndviphi.ts1)



```
plot(predphi.st, cos(modvars.accndvi.phist$accndviphi.ts1), xlab="predicted", ylab="empirical", main="cos")
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 = 4.9932, df = 41, p-value = 1.146e-05
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.3858582 0.7725893
## sample estimates:
##      cor
## 0.614937
```

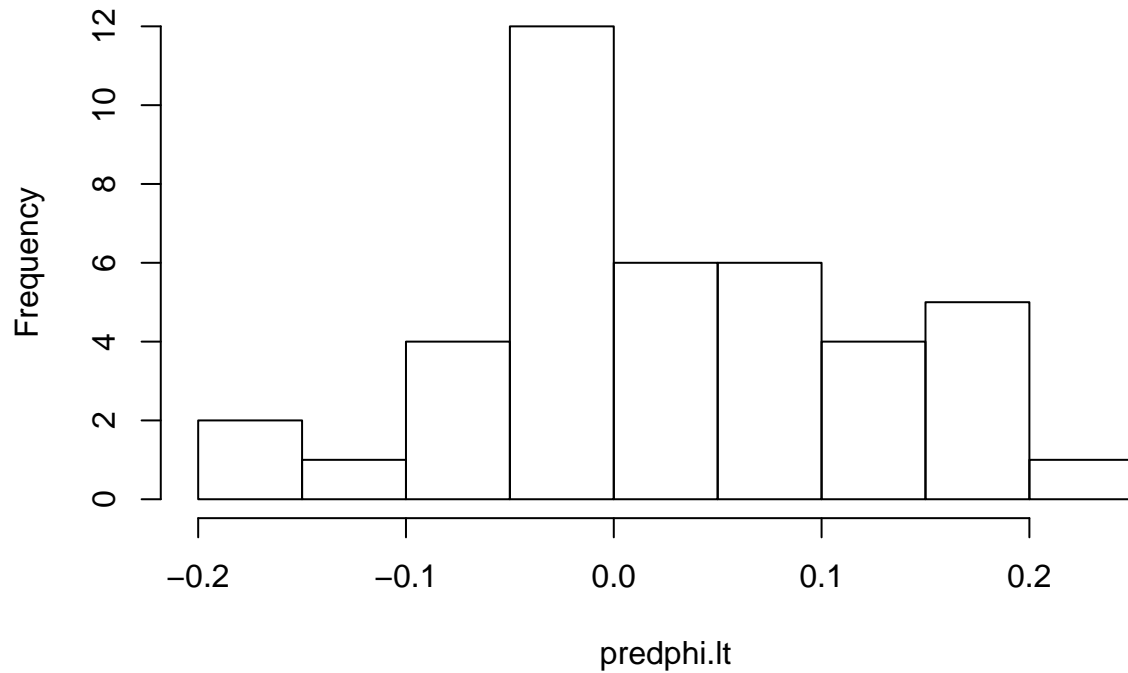
```
varimp.phi.st<-varimp.cforest(cforest.phi.st)
print(varimp.phi.st)
```

```
##      lake_area_ha lake_perim_meters      maxdepth      pct.ag
##      0.14335061      0.10553978      0.17409218      -0.02426418
##           chla           tsi      hu4_zoneid      cv.accndvi
##      -0.06902430      -0.06675996      -0.04438473      -0.02740571
```

```
cforest.phi.lt<-cforest(cos(accndviphi.ts2) ~ lake_area_ha + lake_perim_meters + maxdepth + pct.ag + chla,
                        data=modvars.accndvi.philt)
predphi.lt<-predict.cforest(cforest.phi.lt, newdata=modvars.accndvi.philt)
hist(predphi.lt)
```

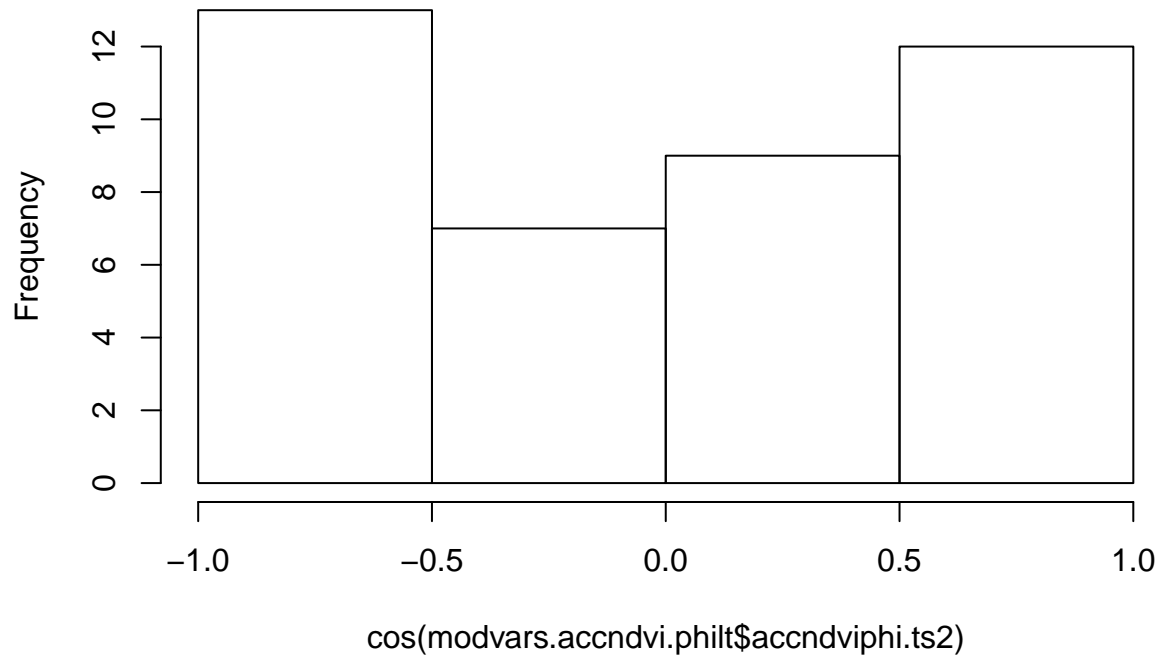


### Histogram of predphi.lt



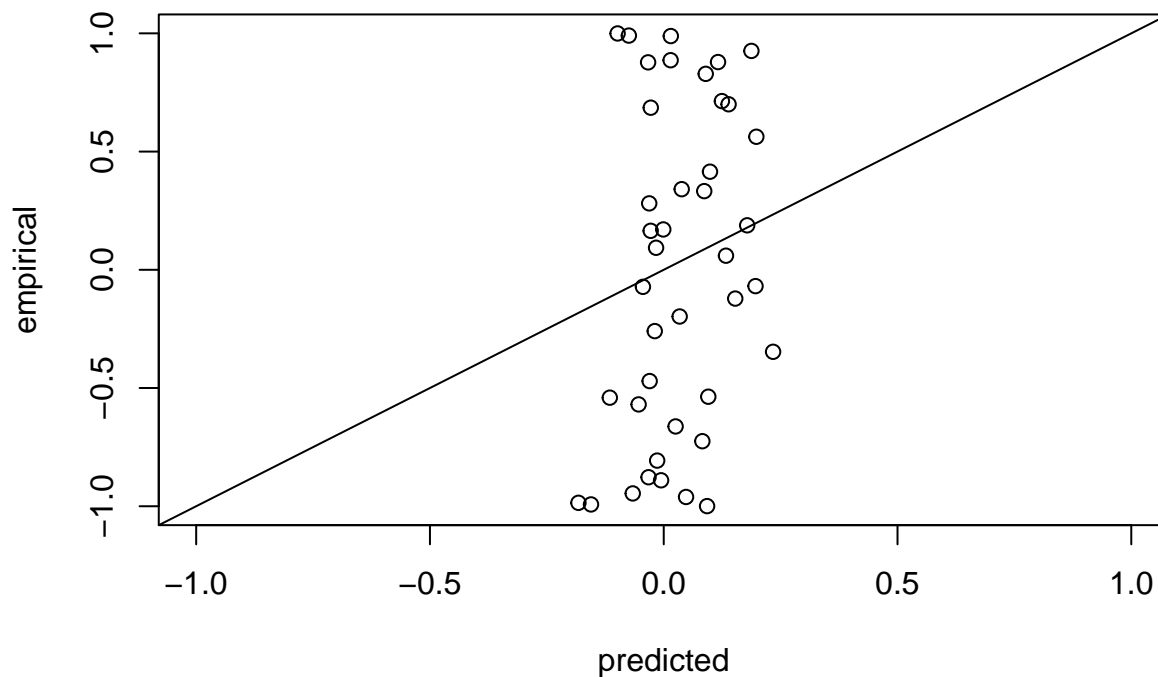
```
hist(cos(modvars.accndvi.philt$accndviphi.ts2))
```

### Histogram of cos(modvars.accndvi.philt\$accndviphi.ts2)



```
plot(predphi.lt, cos(modvars.accndvi.philt$accndviphi.ts2), xlab="predicted", ylab="empirical", main="cosine")
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 = 1.7412, df = 39, p-value = 0.08954
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## -0.04260492 0.53224017
## sample estimates:
## cor
## 0.2685657
```

```
varimp.phi.lt<-varimp.cforest(cforest.phi.lt)
print(varimp.phi.lt)
```

```
##      lake_area_ha lake_perim_meters      maxdepth      pct.ag
##      0.10708894      0.08151034      -0.03946809      -0.04589118
##           chla           tsi      hu4_zoneid      cv.accndvi
##      -0.04451058      -0.12420187      0.15331937      -0.01685269
```