

Homework #3

Cameron Bracken
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Problem 1

```
source('lib.R')

sst <- as.matrix(read.table('data/kaplan-sst-wy-1925-2003-revised.txt'))
sst <- t(matrix(sst[,3],nrow=length(unique(sst[,1])),byrow=T))
sst.lat <- as.matrix(read.table('data/kaplan-sst-wy-1925-2003-II.txt'))
sst.lon <- sst.lat[,2]; sst.lat <- sst.lat[,1]
sst.lon[sst.lon < 0] <- sst.lon[sst.lon < 0] + 360

pdsi <- as.matrix(read.table('data/pdsi-wy-1925-2003.txt'))
pdsi <- t(matrix(pdsi[,3],nrow=length(unique(pdsi[,1])),byrow=T))
pdsi.lat <- as.matrix(read.table('data/pdsi-wy-1925-2003-II.txt'))
pdsi.lon <- pdsi.lat[,2]; pdsi.lat <- pdsi.lat[,1]
pdsi.lon[pdsi.lon < 0] <- pdsi.lon[pdsi.lon < 0] + 360

pacific <- sst.lat > -20 & sst.lon >= 120 & sst.lon <= 280
atlantic <- sst.lat > -20 & sst.lat < 70 & sst.lon >= 250 & sst.lon <= 360
states <- pdsi.lat > 15 & pdsi.lat < 60 & pdsi.lon >= 230 & pdsi.lon <= 295

lon.pac <- sst.lon[pacific]
lat.pac <- sst.lat[pacific]
sst.pac <- sst[,pacific]
lon.atl <- sst.lon[atlantic]
lat.atl <- sst.lat[atlantic]
sst.atl <- sst[,atlantic]
lon.usa <- pdsi.lon[states]
lat.usa <- pdsi.lat[states]
pdsi.usa <- pdsi[,states]

glo <- my.pca(sst)
pac <- my.pca(sst.pac)
atl <- my.pca(sst.atl)
usa <- my.pca(pdsi.usa)

save(lat, lon, lat.pac, lon.pac, lat.atl, lon.atl, lat.usa, lon.usa, pac, atl,
      glo, usa, sst, sst.lat, sst.lon, sst.pac, sst.atl, pdsi.usa,
      file='output/1.Rdata')
```

Figure 1: Reading the data and calculating the statistics.

```
my.pca <- function(x){  
  s <- svd(var(x))  
  
  #do an Eigen decomposition..  
  eigv <- s$u  
  eig <- s$d  
  
  #Principal Components..  
  pcs <- t(t(eigv) %*% t(x))  
  
  #Eigen Values.. - fraction variance  
  eigf <- (eig/sum(eig))  
  
  return(list(eigv=eigv,eigf=eigf,eig=eig,pc=pcs))  
}
```

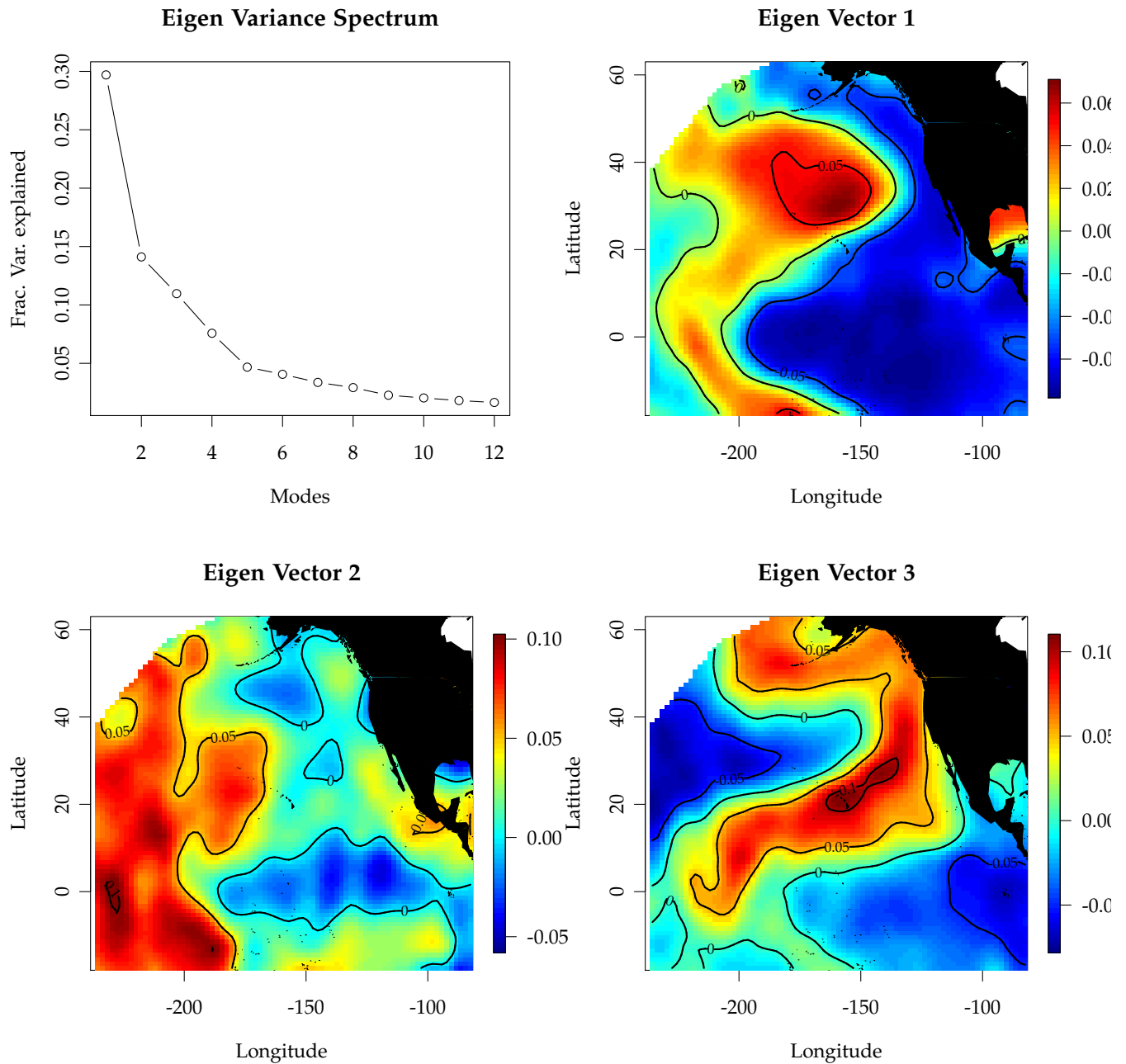
Figure 2: PCA function.

Problem 1 (i)

```

> layout(rbind(c(1,2),c(3,4)))
> plot(pac$eigf[1:12], type="b", xlab="Modes",
+       ylab="Frac. Var. explained",main="Eigen Variance Spectrum")
> for(i in 1:3){
+   surf <- Tps(cbind(lon.pac-360,lat.pac),pac$eigv[,i])
+   surface(surf,xlab="Longitude", ylab="Latitude",main=sprintf("Eigen Vector %d",i))
+   map('world',add=TRUE,fill=TRUE)
+ }

```

**Figure 3:** Pacific SST

```

> layout(cbind(c(1,2,3)))
> for(i in 1:3){
+   plot(ts(pac$pc[,i],start=1925,frequency=1),
+         main=sprintf("PC %d",i),ylab='')
+ }

```

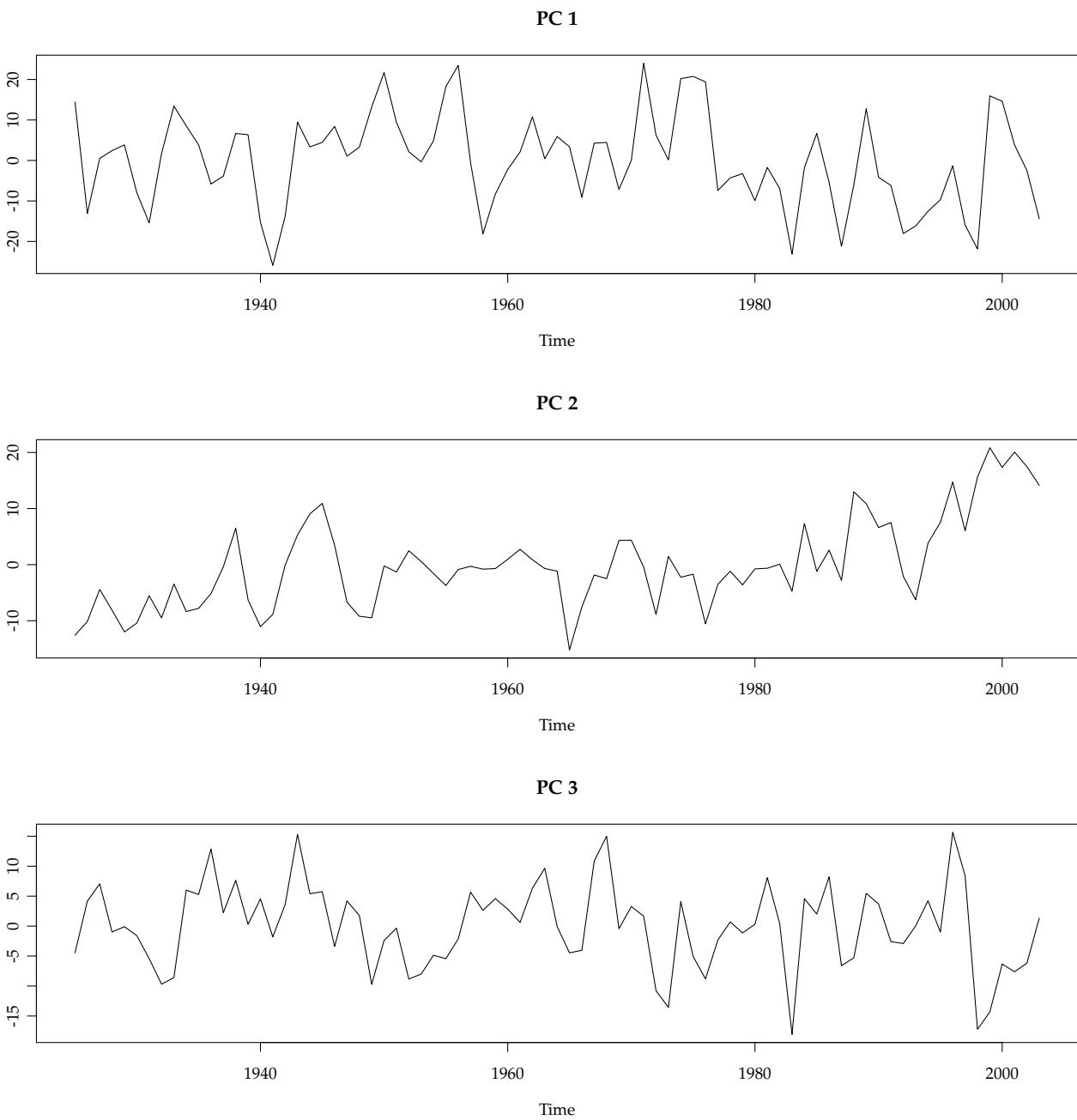


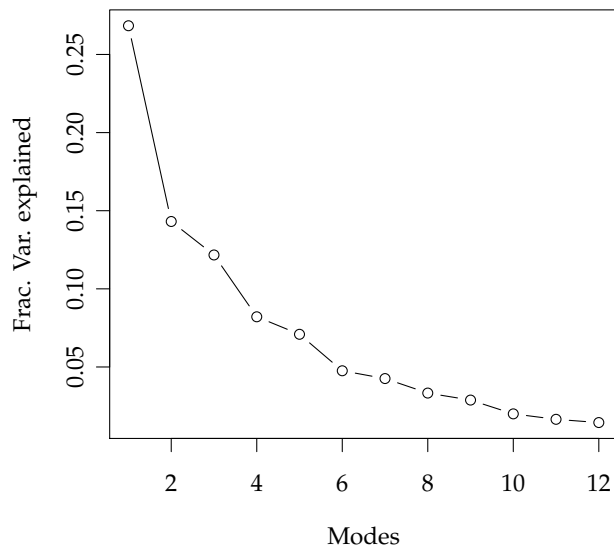
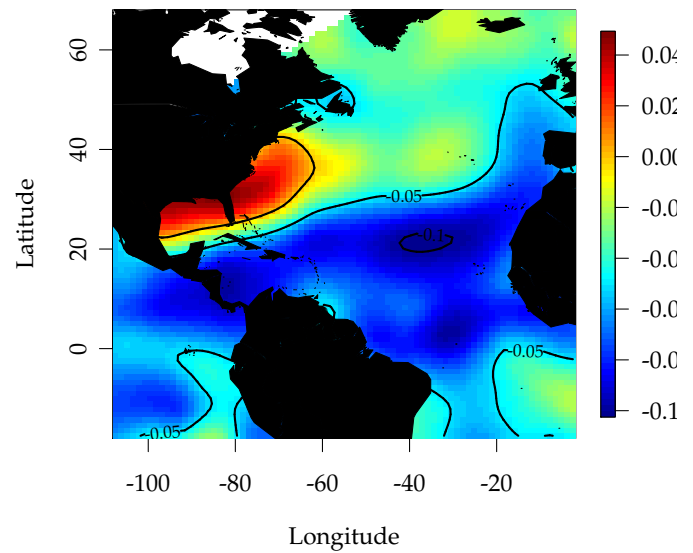
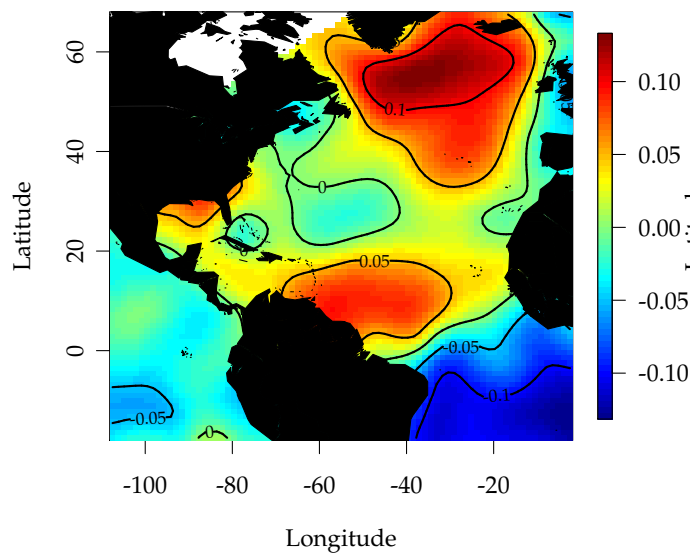
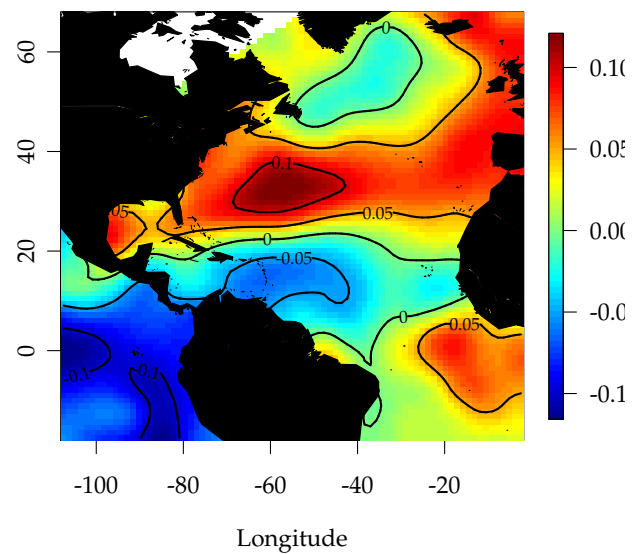
Figure 4: PC's of Pacific SST

Problem 1 (ii)

```

> layout(rbind(c(1,2),c(3,4)))
> plot(atl$eigf[1:12], type="b", xlab="Modes",
+       ylab="Frac. Var. explained",main="Eigen Variance Spectrum")
> for(i in 1:3){
+   surf <- Tps(cbind(lon.atl-360,lat.atl),atl$eigv[,i])
+   surface(surf,xlab="Longitude", ylab="Latitude",main=sprintf("Eigen Vector %d",i))
+   map('world',add=TRUE,fill=T)
+ }

```

Eigen Variance Spectrum**Eigen Vector 1****Eigen Vector 2****Eigen Vector 3****Figure 5: Atlantic SST**

```

> layout(cbind(c(1,2,3)))
> for(i in 1:3){
+   plot(ts(atl$pc[,i],start=1925,frequency=1),
+         main=sprintf("PC %d",i),ylab='')
+ }

```

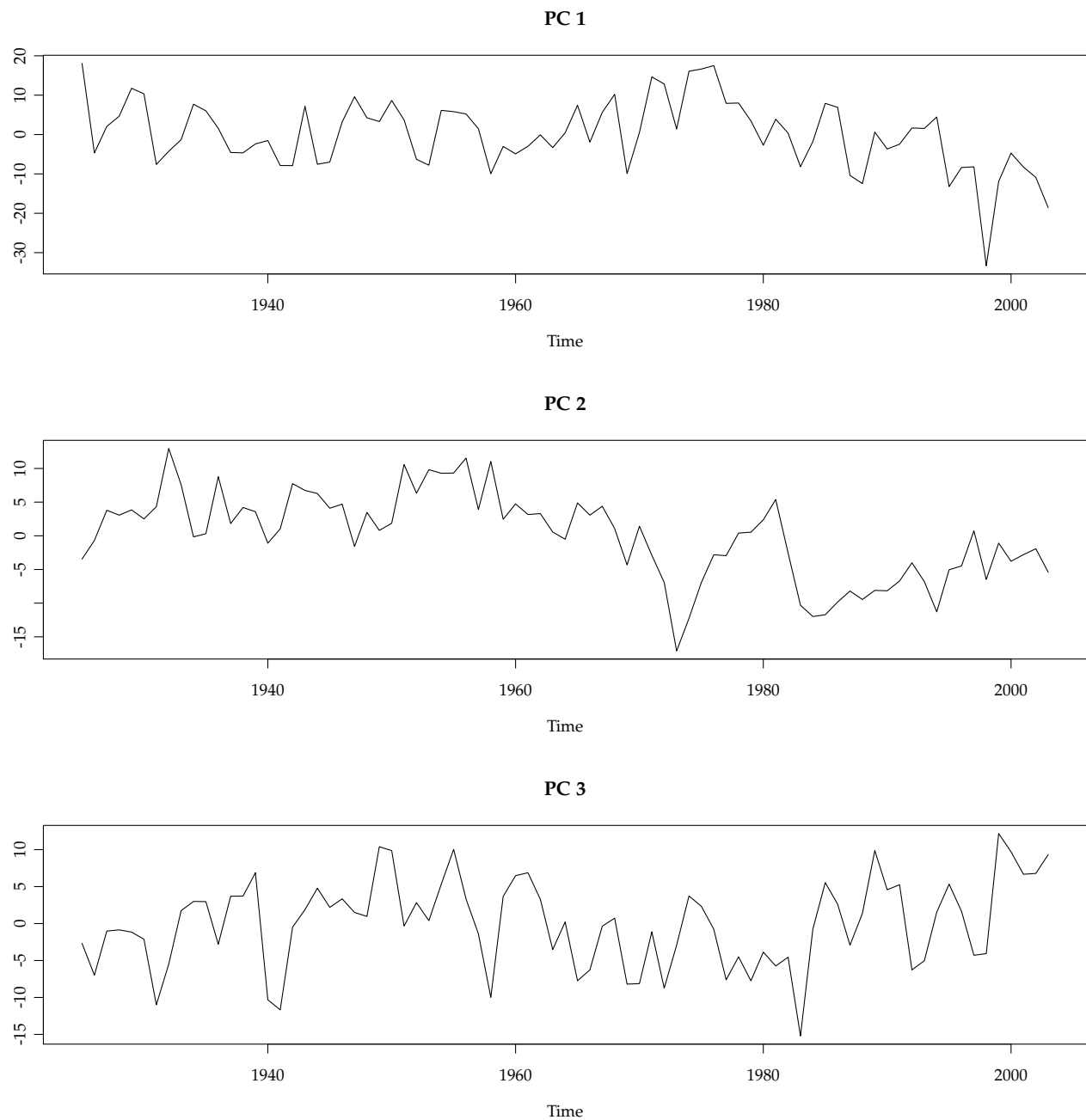


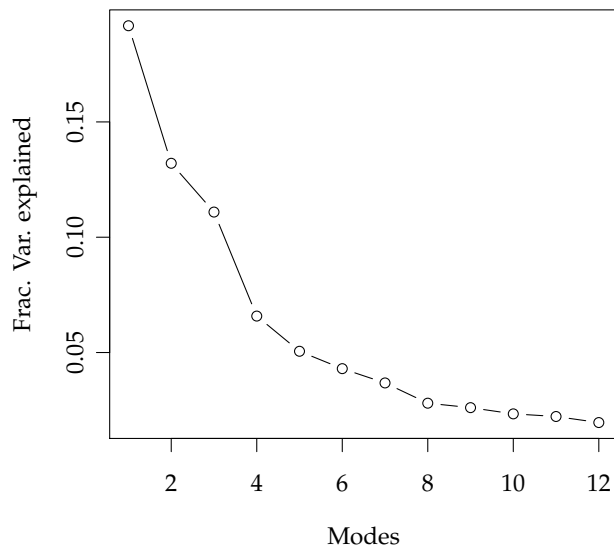
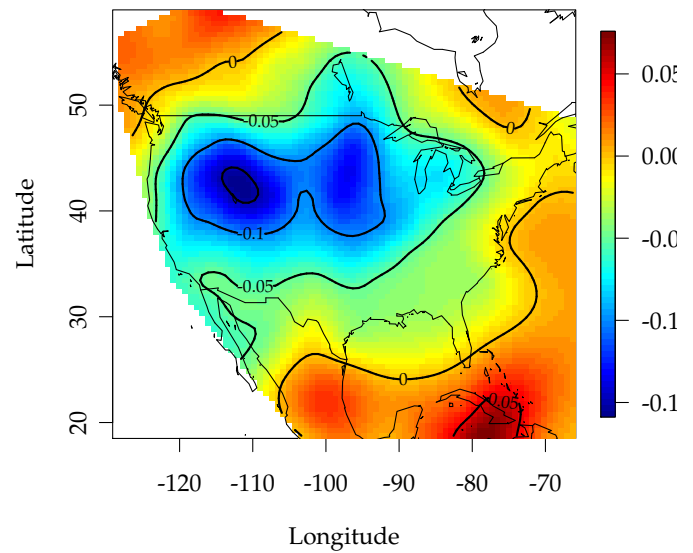
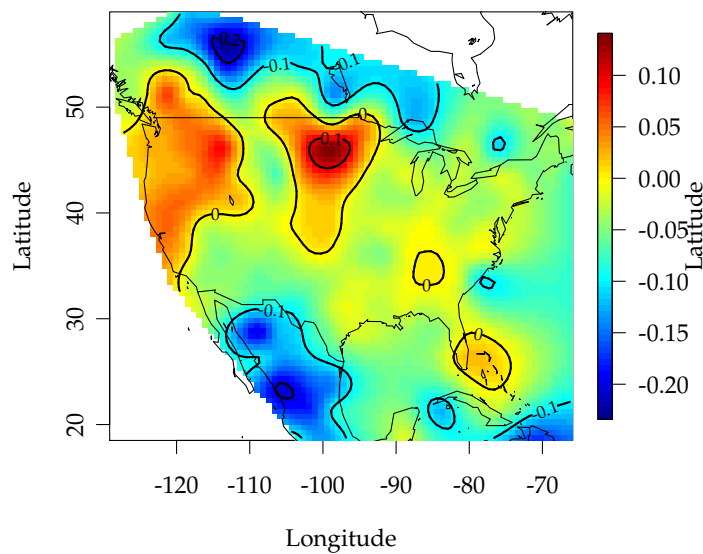
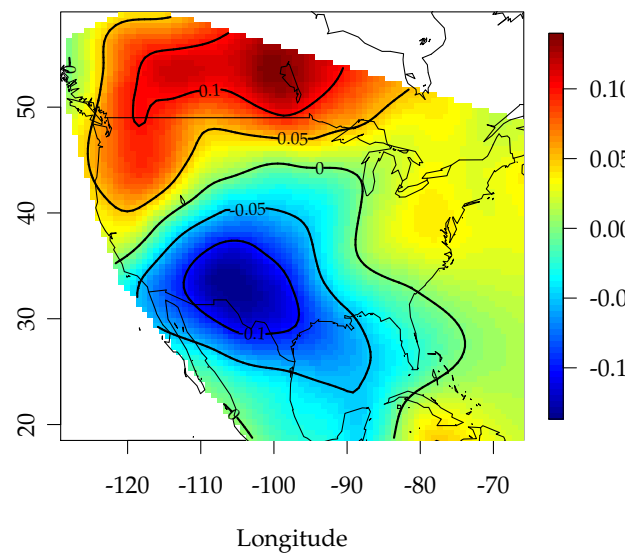
Figure 6: PC's of Atlantic SST

Problem 1 (iii)

```

> layout(rbind(c(1,2),c(3,4)))
> plot(usa$eigf[1:12], type="b", xlab="Modes",
+       ylab="Frac. Var. explained",main="Eigen Variance Spectrum")
> for(i in 1:3){
+   surf <- Tps(cbind(lon.usa-360,lat.usa),usa$eigv[,i])
+   surface(surf,xlab="Longitude", ylab="Latitude",main=sprintf("Eigen Vector %d",i))
+   map('world',add=TRUE)
+ }

```

Eigen Variance Spectrum**Eigen Vector 1****Eigen Vector 2****Eigen Vector 3****Figure 7: PDSI over the United States**

```

> layout(cbind(c(1,2,3)))
> for(i in 1:3){
+   plot(ts(usa$pc[,i],start=1925,frequency=1),
+         main=sprintf("PC%d",i),ylab='')
+ }

```

PC1



PC2



PC3



Figure 8: PC's of US PDSI

Problem 2

Below is the code to calculate the correlation maps and local polynomial fits for SST and PDSI.

```
source('lib.R')
if(file.exists('output/1.Rdata')) load('output/1.Rdata') else source('1.R')
library(locfit)

#usa - x
#pac - y
lfpc <- list()
for(i in 1:2){
  for(j in 1:2){
    b <- best.par(pac$pc[,i],usa$pc[,j])
    lfpc[[paste(i,j,sep=' ')] <- locfit(usa$pc[,j]~pac$pc[,i],alpha=b$a,deg=b$p)
  }
}

#correlation maps
cor.pdsi.sst.pac1 <- as.vector(cor(pac$pc[,1],pdsi.usa))
cor.pdsi.sst.pac2 <- as.vector(cor(pac$pc[,2],pdsi.usa))
cor.pdsi.sst.atl1 <- as.vector(cor(atl$pc[,1],pdsi.usa))
cor.pdsi.sst.atl2 <- as.vector(cor(atl$pc[,2],pdsi.usa))

cor.map <- list()
cor.map[[1]] <- Tps(cbind(lon.usa-360,lat.usa),cor.pdsi.sst.pac1)
cor.map[[2]] <- Tps(cbind(lon.usa-360,lat.usa),cor.pdsi.sst.pac2)
cor.map[[3]] <- Tps(cbind(lon.usa-360,lat.usa),cor.pdsi.sst.atl1)
cor.map[[4]] <- Tps(cbind(lon.usa-360,lat.usa),cor.pdsi.sst.atl2)

save(lfpc,cor.map,file='output/2.Rdata')
```

Figure 9: Code for problem 2.

```

> layout(rbind(c(1,2),c(3,4)))
> for(i in 1:2)
+   for(j in 1:2){
+     plot(pac$pc[,i],usa$pc[,j],
+         xlab=sprintf('sst pc %d',i),
+         ylab=sprintf('pdsi pc %d',j))
+     lines(lfpc[[paste(i,j,sep='')]])
+   }

```

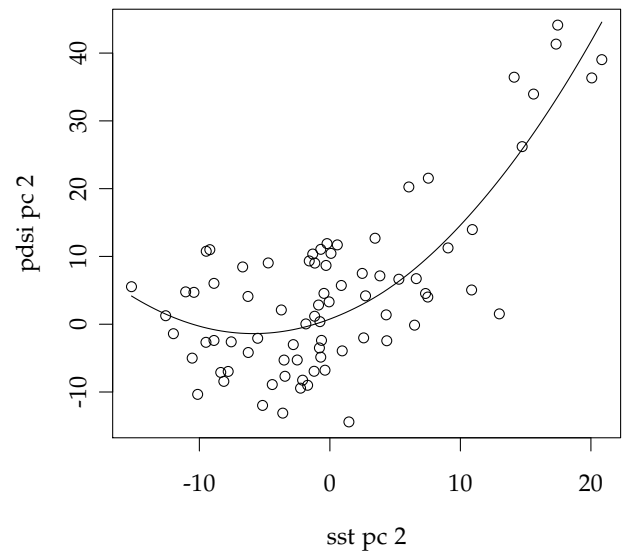
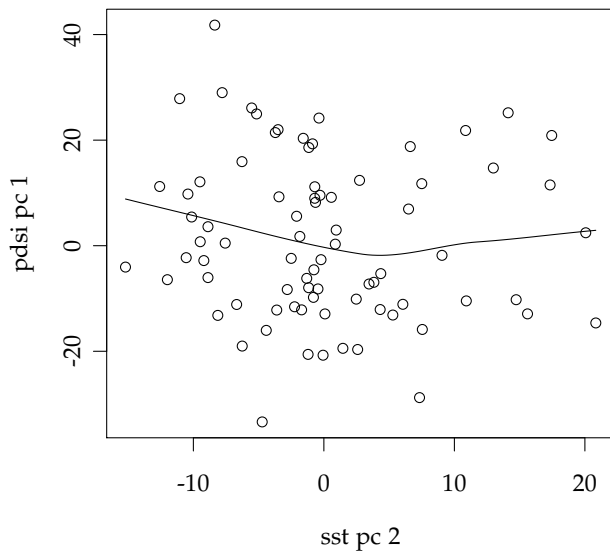
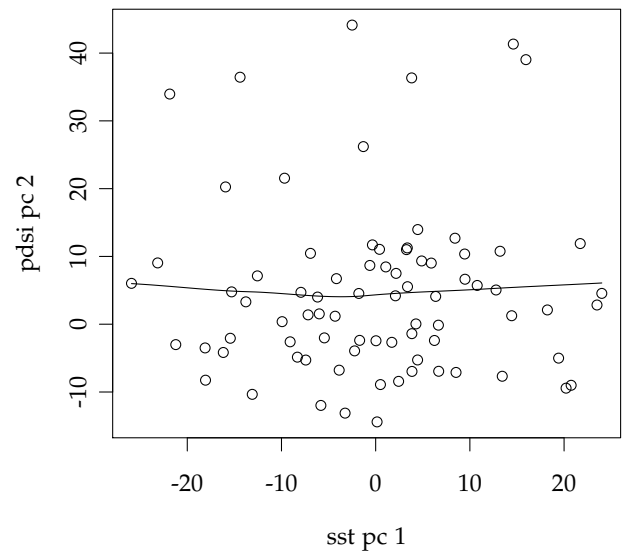
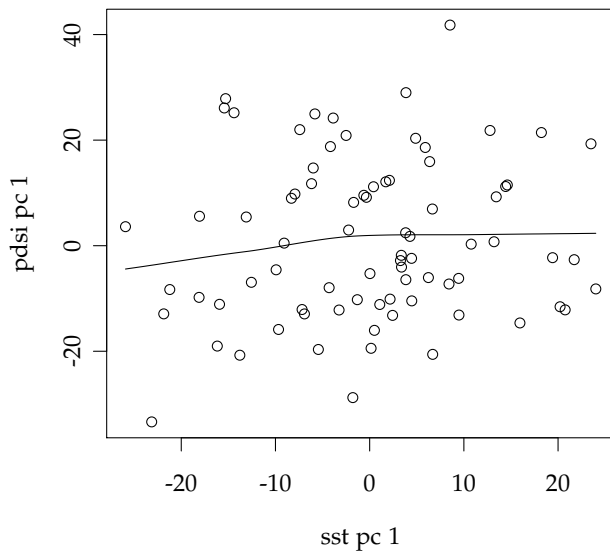


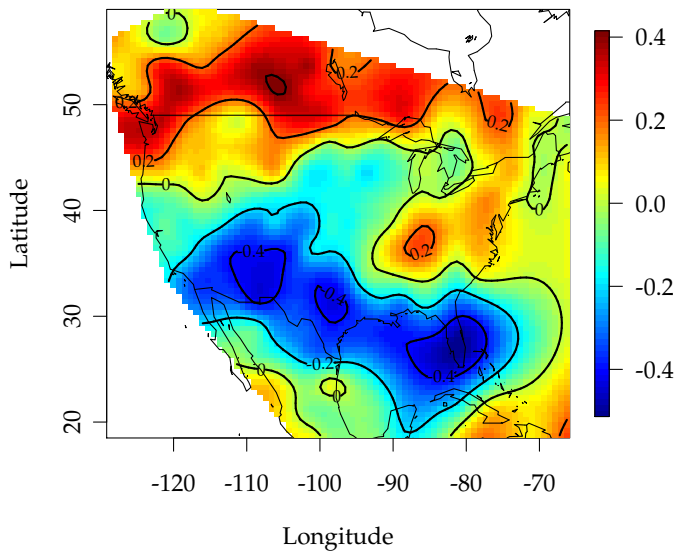
Figure 10: States PC

```

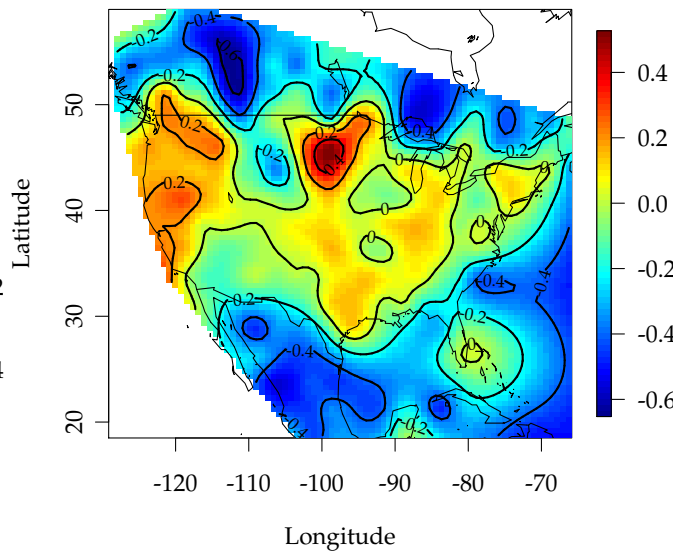
> layout(rbind(c(1,2),c(3,4)))
> for(i in 1:4){
+   surface(cor.map[[i]], xlab="Longitude", ylab="Latitude",
+           main=paste("PDSI correlated with ",
+                       ifelse(i>2,"Atlantic","Pacific")," SST PC",
+                       ifelse(i%%2==0,2,1)))
+   map('world',add=TRUE)
+ }

```

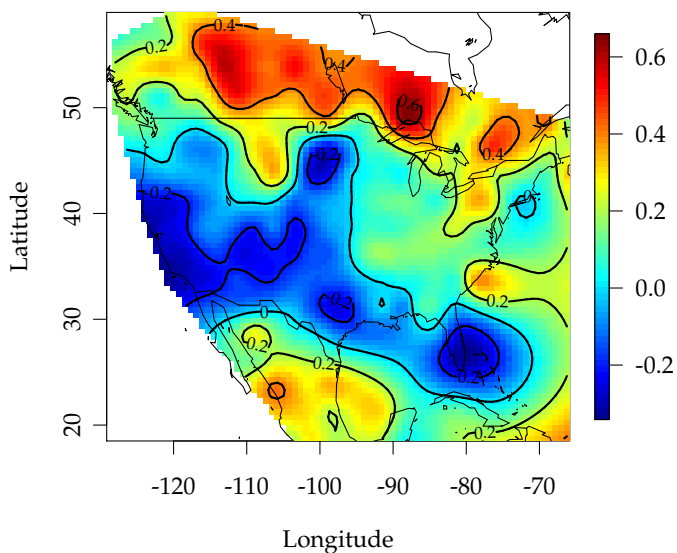
PDSI correlated with Pacific SST PC 1



PDSI correlated with Pacific SST PC 2



PDSI correlated with Atlantic SST PC 1



PDSI correlated with Atlantic SST PC 2

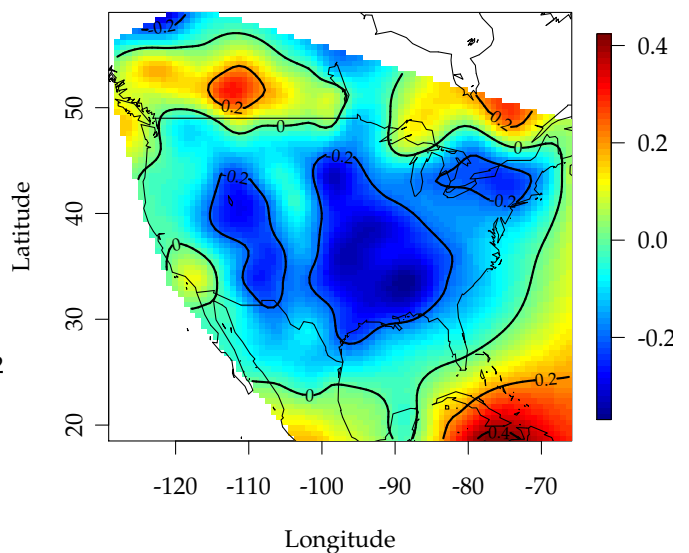


Figure 11: SST principal components correlated with PDSI

Problem 3

Below is the code which performs SVD and calculates the heterogeneous correlation maps.

```
source('lib.R')
if(file.exists('output/1.Rdata')) load('output/1.Rdata') else source('1.R')

C <- var(pdsi.usa,sst)
S <- svd(C)

#Time coefficients
pdsi.tc <- pdsi.usa %*% S$u
sst.tc <- sst %*% S$v

cor.map.sst <- cor.map.pdsi <- list()
for(i in 1:3){
  cor.map.sst[[i]] <-
    Tps(cbind(sst.lon-180,sst.lat),as.vector(cor(sst.tc[,1],sst)))
  cor.map.pdsi[[i]] <-
    Tps(cbind(lon.usa-360,lat.usa),as.vector(cor(pdsi.tc[,1],pdsi.usa)))
}

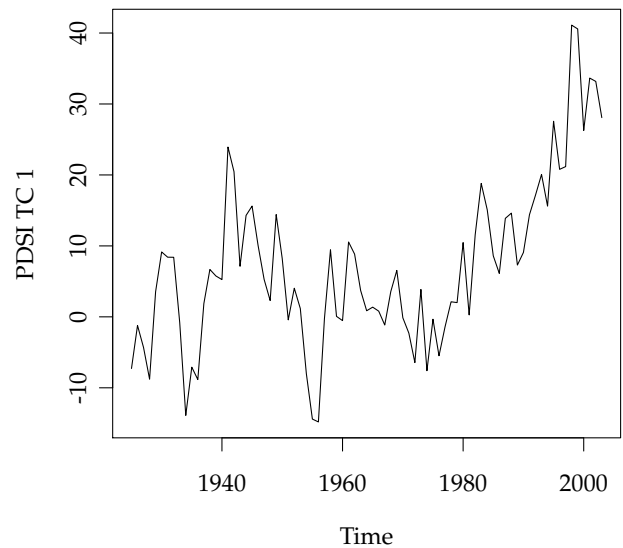
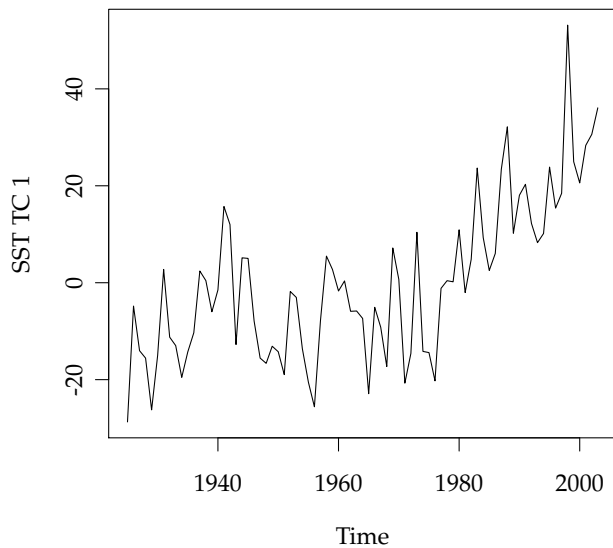
save(sst.tc, pdsi.tc, cor.map.sst, cor.map.pdsi, file='output/3.Rdata')
```

Figure 12: Code for problem 3.

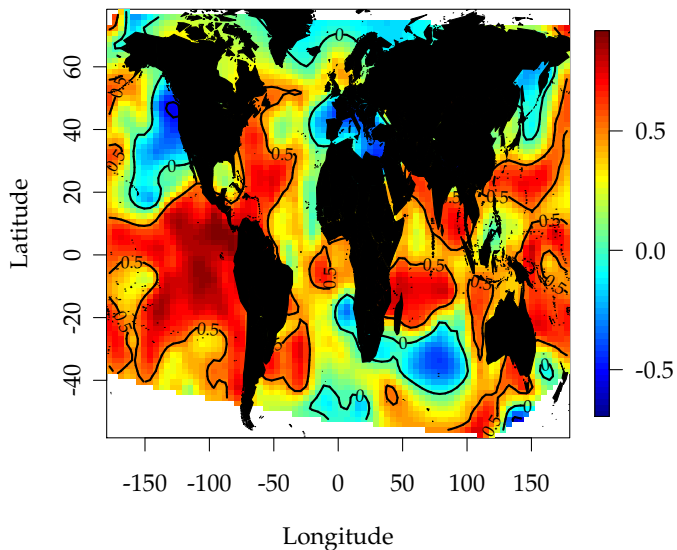
```

> layout(rbind(c(1,2),c(3,4)))
> plot(ts(sst.tc[,1],start=1925,frequency=1),ylab="SST TC 1")
> plot(ts(pdsi.tc[,1],start=1925,frequency=1),ylab="PDSI TC 1")
> surface(cor.map.sst[[1]], xlab="Longitude", ylab="Latitude",
+         main='Het. Correlation map of SST (mode 1)')
> map('world',add=TRUE,fill=TRUE)
> surface(cor.map.pdsi[[1]], xlab="Longitude", ylab="Latitude",
+         main='Het. Correlation map of PDSI (mode 1)')
> map('world',add=TRUE)

```



Het. Correlation map of SST (mode 1)



Het. Correlation map of PDSI (mode 1)

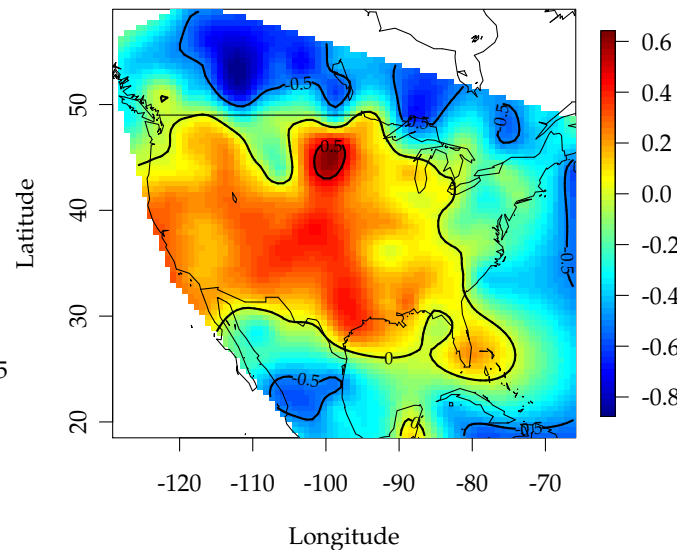
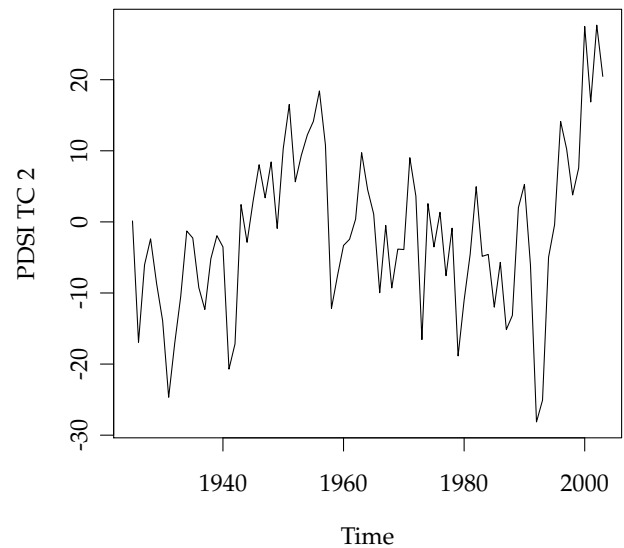
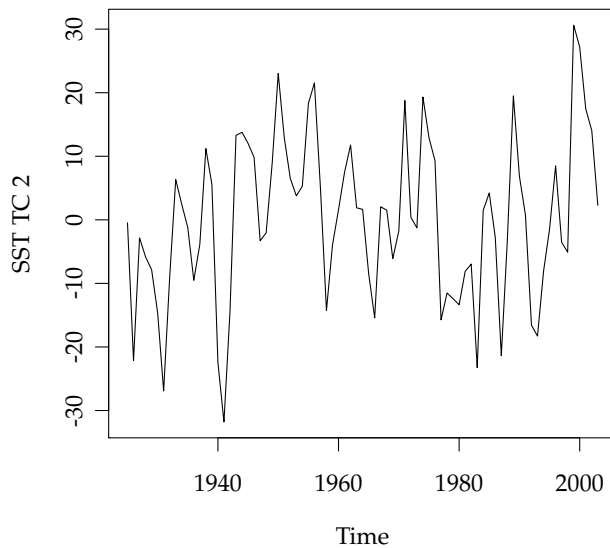


Figure 13: Heterogeneous correlation maps and time coefficients for mode 1

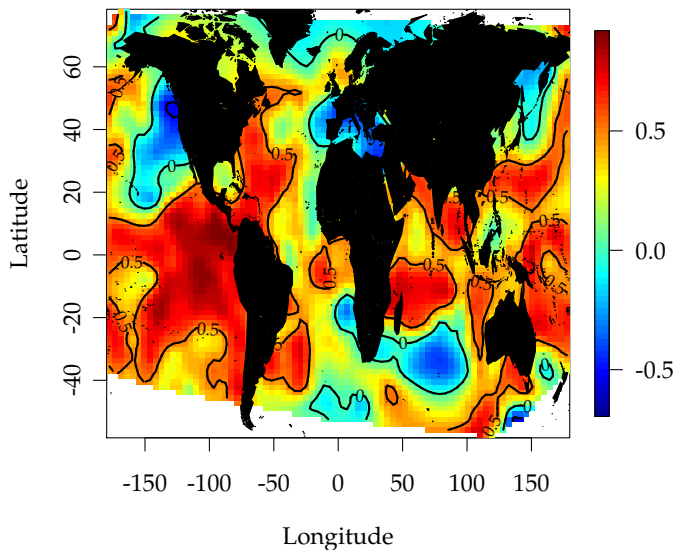
```

> layout(rbind(c(1,2),c(3,4)))
> plot(ts(sst.tc[,2],start=1925,frequency=1),ylab="SST TC 2")
> plot(ts(pdsi.tc[,2],start=1925,frequency=1),ylab="PDSI TC 2")
> surface(cor.map.sst[[2]], xlab="Longitude", ylab="Latitude",
+         main='Het. Correlation map of SST (mode 2)')
> map('world',add=TRUE,fill=TRUE)
> surface(cor.map.pdsi[[2]], xlab="Longitude", ylab="Latitude",
+         main='Het. Correlation map of PDSI (mode 2)')
> map('world',add=TRUE)

```



Het. Correlation map of SST (mode 2)



Het. Correlation map of PDSI (mode 2)

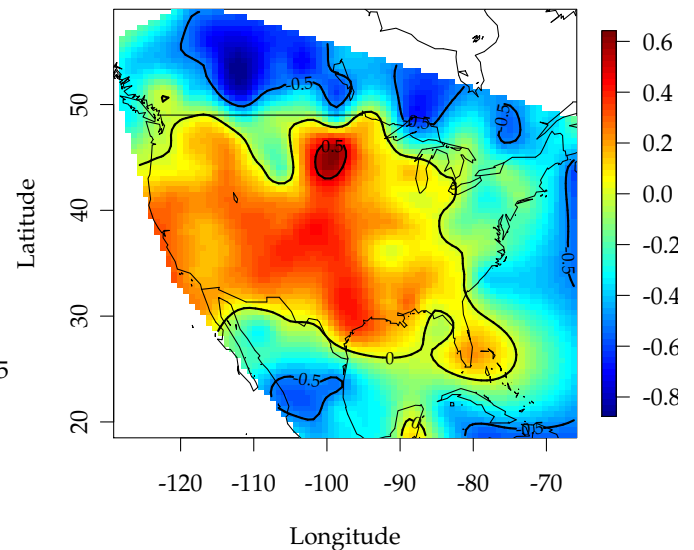
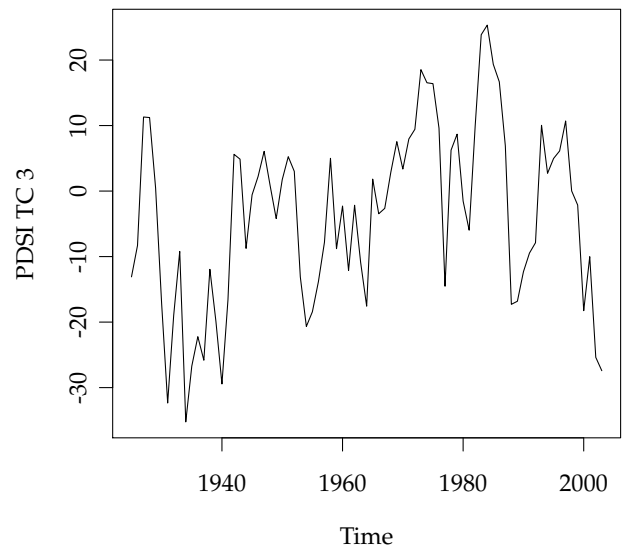
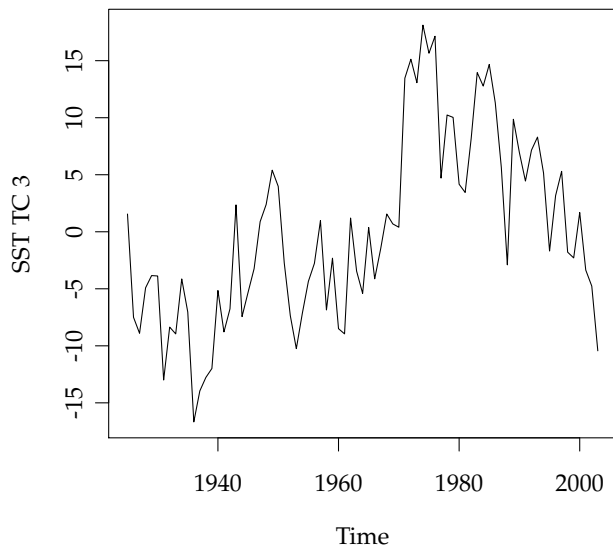


Figure 14: Heterogeneous correlation maps and time coefficients for mode 2

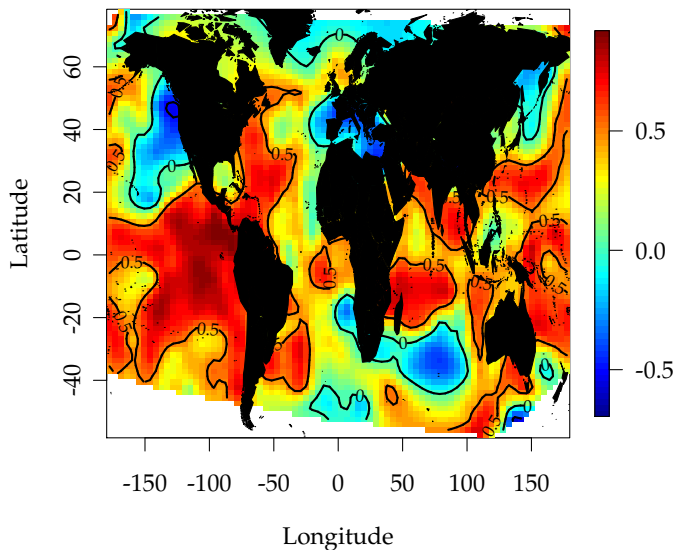
```

> layout(rbind(c(1,2),c(3,4)))
> plot(ts(sst.tc[,3],start=1925,frequency=1),ylab="SST TC 3")
> plot(ts(pdsi.tc[,3],start=1925,frequency=1),ylab="PDSI TC 3")
> surface(cor.map.sst[[3]], xlab="Longitude", ylab="Latitude",
+         main='Het. Correlation map of SST (mode 3)')
> map('world',add=TRUE,fill=TRUE)
> surface(cor.map.pdsi[[3]], xlab="Longitude", ylab="Latitude",
+         main='Het. Correlation map of PDSI (mode 3)')
> map('world',add=TRUE)

```



Het. Correlation map of SST (mode 3)



Het. Correlation map of PDSI (mode 3)

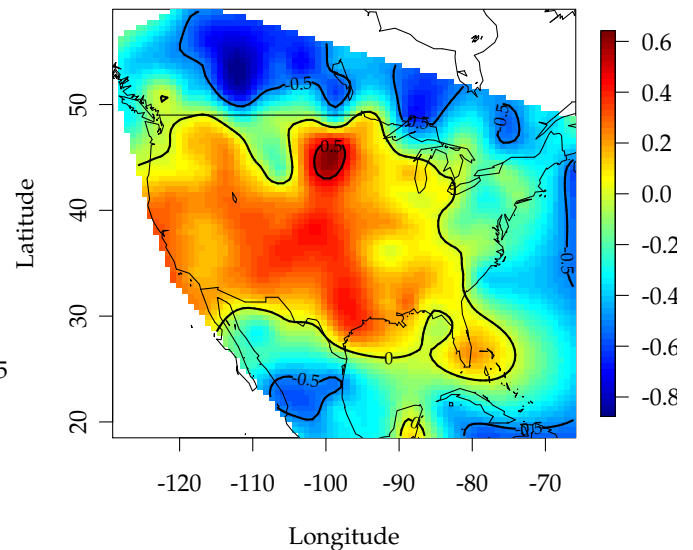


Figure 15: Heterogeneous correlation maps and time coefficients for mode 3

Problem 4

```

source('lib.R')
flow <- as.matrix(read.table("data/AMJJFLOW-49-99.txt"))/(10^3)
swe <- as.matrix(read.table("data/March1SWE-1949-99.txt"))
nsim <- 250

#Singular value decomposition
C <- var(flow,swe) #note that the variable with fewer columns should be first
S <- svd(C)

#Time coefficients
flow.tc <- flow %*% S$u
swe.tc <- swe %*% S$v

#heterogeneous Correlation maps
cor.map.swe <- cor.map.flow <- list()
for(i in 1:3){
  cor.map.swe[[i]] <- as.vector(cor(swe.tc[,1],swe))
  cor.map.flow[[i]] <- as.vector(cor(flow.tc[,1],flow))
}

#Local polynomial fit of TC
lftc <- list()
for(i in 1:2)
  for(j in 1:2){
    b <- best.par(swe.tc[,i],flow.tc[,j])
    lftc[[paste(i,j,sep=' ')] <-
      locfit(flow.tc[,j]~swe.tc[,i],alpha=b$a,deg=b$p)
  }

#Forecasting
ypred.svd <- fc.svd(flow,swe,nsim)

#get the rpss and median correlation stats
stats.svd <- fc.stats(flow,ypred.svd,nsim)
stats.svd$rpss <-

save(swe.tc, flow.tc, cor.map.swe, cor.map.flow, lftc, stats.svd, ypred.svd,
     flow, file='output/4.Rdata')

```

Figure 16: SVD code.


```

fc.svd <- function(flow,swe,nsim){
  n <- nrow(flow)
  n.sites <- ncol(flow)
  nsim <- 100
  ypred <- array(NA,c(n.sites,nsim,n))
  yy <- numeric(n.sites)

  for(i in 1:n){
    #drop a point..
    swe.cv <- swe[-i,]
    flow.cv <- flow[-i,]

    #perform SVD
    C <- var(flow.cv,swe.cv)
    S <- svd(C)

    #Get the tc's
    flow.cv.tc <- flow.cv %%% S$u
    swe.cv.tc <- swe.cv %%% S$v
    cv.model <- lsfit(swe.cv.tc[,1],flow.cv.tc[,1])
    stderr <- sum((cv.model$resid)^2) / (nrow(flow.cv) - 2)

    #prediction error..
    Sxx <- sum((swe.cv.tc[,1] - mean(swe.cv.tc[,1]))^2)

    #the TCs of the dropped point..
    xp <- swe[i,] %%% S$v

    #generate an ensemble..
    for(isim in 1:nsim){
      this.stderr <- sqrt(stderr *
        (1 + (1/(n-1)) +
          ((xp[1] - mean(swe.cv.tc[,1]))^2)/Sxx))

      yy[1] <- cv.model$coef[1] +
        cv.model$coef[2]*xp[1] + rnorm(1,0,this.stderr)

      #grab random TCs for the remaining TCs
      for(site in 2:n.sites){

        r.tc <- round(runif(1,1,(n-1)))
        yy[site] <- flow.cv.tc[r.tc,site]
      }
      ypred[1:n.sites,isim,i]=yy %%% t(S$u)
    }
  }
  ypred[ypred < 0] <- 0
  return(ypred)
}

```

Figure 17: SVD forecast code.

```

fc.stats <- function(flow,ypred,nsim){

  n.sites <- ncol(flow)
  site.rpss <- site.mc <- numeric(n.sites)
  n <- nrow(flow)

  ##Calculate rpss
  for(i in 1:n.sites){

    thresh <- quantile(flow[,i], c(0.33, 0.66))

    # climatological forecast..
    climo=cumsum(c(1/3, 1/3, 1/3))

    rpss = numeric(n)
    for(jj in 1:n){

      # forecast categorical probabilities
      fcastprob=1:3
      yypred=ypred[i,,jj]
      fcastprob[1] <- length(yypred[yypred <= thresh[1]]) / nsim
      fcastprob[2] <-
        length(yypred[yypred > thresh[1] &
          yypred < thresh[2]]) / nsim
      fcastprob[3] <- length(yypred[yypred >= thresh[2]]) / nsim

      fcastprob=cumsum(fcastprob)

      # actual..
      actual=rep(0,3)
      if(flow[jj,i] <= thresh[1])actual[1]=1
      if(flow[jj,i] > thresh[1] & flow[jj,i] < thresh[2])actual[2]=1
      if(flow[jj,i] >= thresh[2])actual[3]=1

      rpsclimo = sum((climo-actual)^2)
      rpsfcast = sum((fcastprob - actual)^2)
      rpss[jj] = 1 - (rpsfcast/rpsclimo)

    }
    #print out the median RPSS
    site.rpss[i] <- median(rpss)
    site.mc[i] <- cor(flow[,i],apply(ypred[i,,],2,median))
  }
  return(list(rpss=site.rpss,mc=site.mc))
}

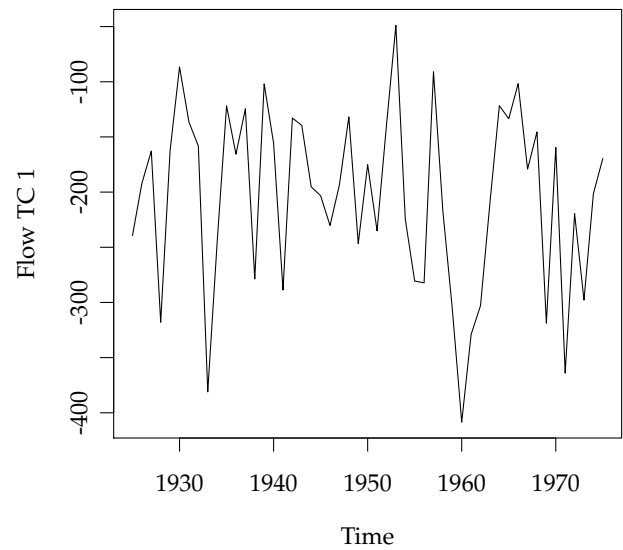
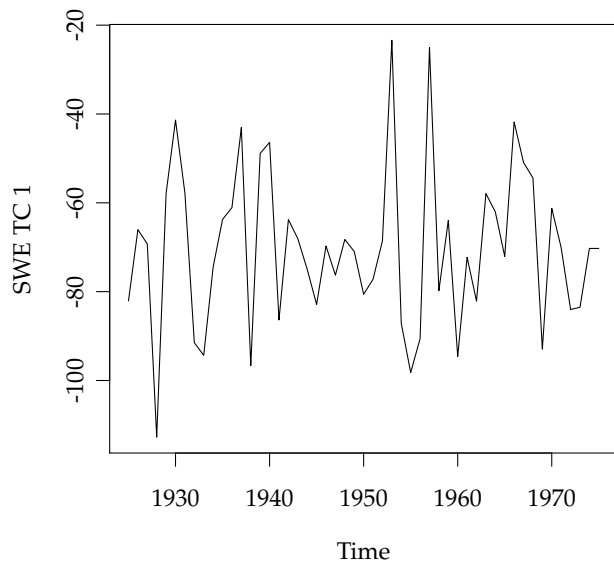
```

Figure 18: SVD forecast stats code.

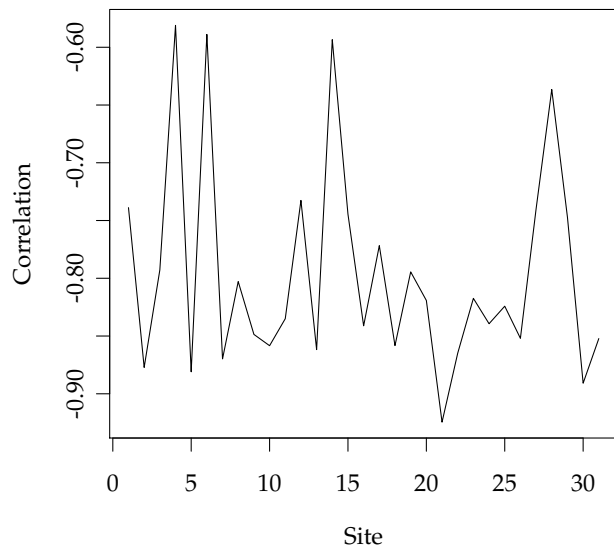
```

> layout(rbind(c(1,2),c(3,4)))
> plot(ts(swe.tc[,1],start=1925,frequency=1),ylab="SWE TC 1")
> plot(ts(flow.tc[,1],start=1925,frequency=1),ylab="Flow TC 1")
> plot(cor.map.swe[[1]], xlab="Site", ylab="Correlation",
+       main='Het. Correlation map of SWE (mode 1)',type='l')
> plot(cor.map.flow[[1]], xlab="Site", ylab="Correlation",
+       main='Het. Correlation map of Flow (mode 1)',type='l')

```



Het. Correlation map of SWE (mode 1)



Het. Correlation map of Flow (mode 1)

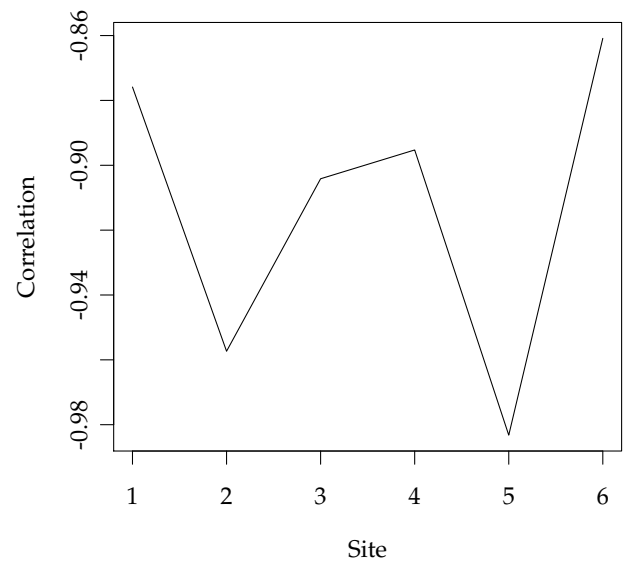
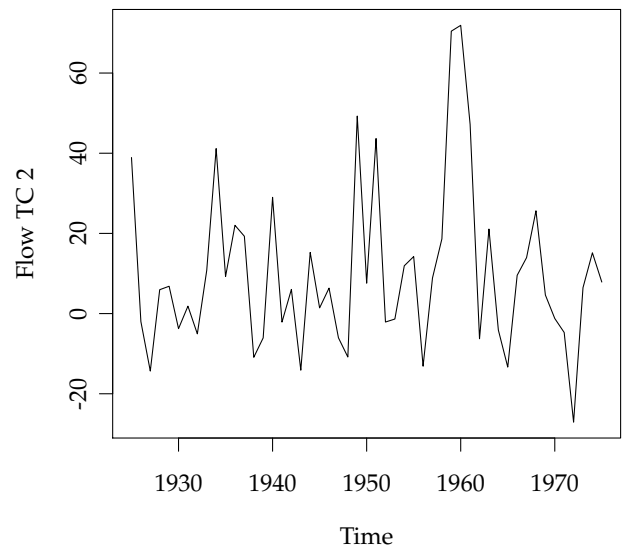
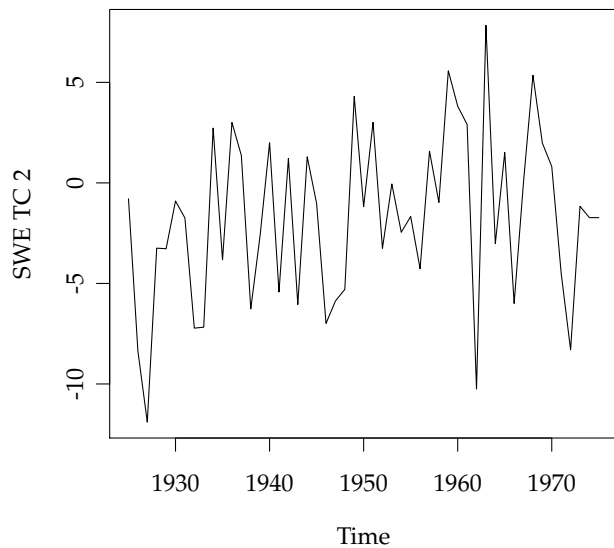


Figure 19: Heterogeneous correlation maps and time coefficients for mode 1

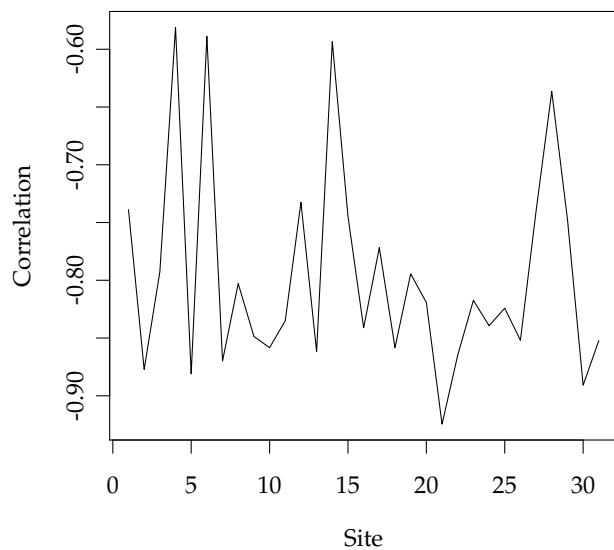
```

> layout(rbind(c(1,2),c(3,4)))
> plot(ts(swe.tc[,2],start=1925,frequency=1),ylab="SWE TC 2")
> plot(ts(flow.tc[,2],start=1925,frequency=1),ylab="Flow TC 2")
> plot(cor.map.swe[[2]], xlab="Site", ylab="Correlation",
+      main='Het. Correlation map of SWE (mode 2)',type='l')
> plot(cor.map.flow[[2]], xlab="Site", ylab="Correlation",
+      main='Het. Correlation map of Flow (mode 2)',type='l')

```



Het. Correlation map of SWE (mode 2)



Het. Correlation map of Flow (mode 2)

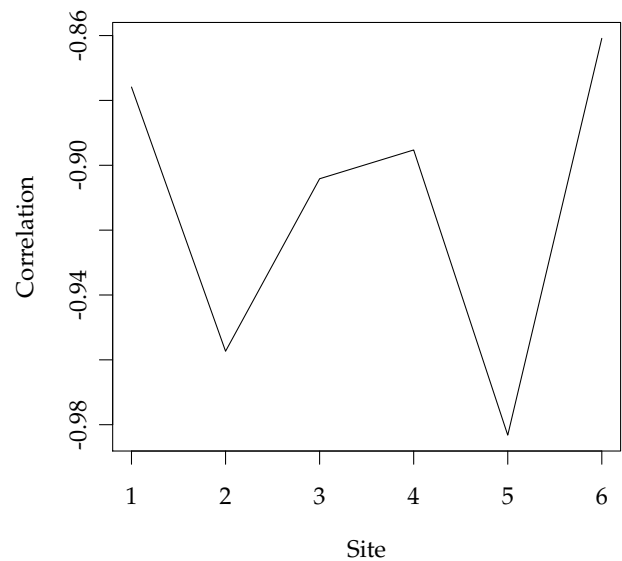


Figure 20: Heterogeneous correlation maps and time coefficients for mode 2

```

> layout(rbind(c(1,2),c(3,4)))
> for(i in 1:2)
+   for(j in 1:2){
+     plot(swe.tc[,i],flow.tc[,j],
+         xlab=sprintf('SWE TC %d',i),
+         ylab=sprintf('Flow TC %d',j))
+     lines(lftc[[paste(i,j,sep='')]])
+   }

```

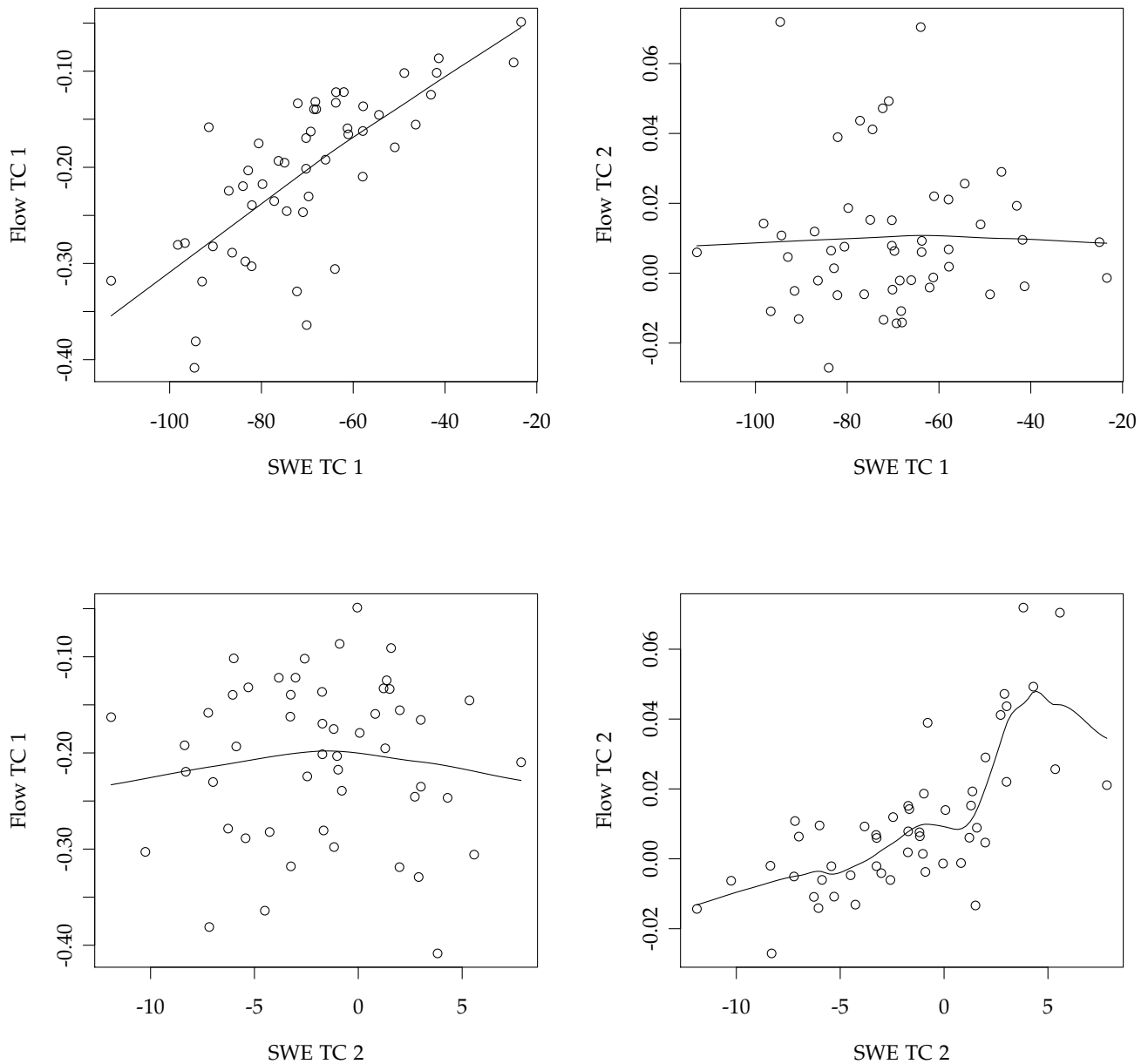


Figure 21: States PC

```

> layout(cbind(c(1:3)))
> for(i in 1:3){
+   this.site <- ypred.svd[i,,]
+   this.site <- as.data.frame(this.site)
+   names(this.site) <- 1949:1999
+   boxplot(this.site,xlab=paste('Site',i))
+   lines(flow[,i],col='blue')
+   mtext(paste('RPSS = ',round(stats.svd$rpss[i],3),
+   'MC = ',round(stats.svd$rpss[i],3)),line=1)
+ }

```

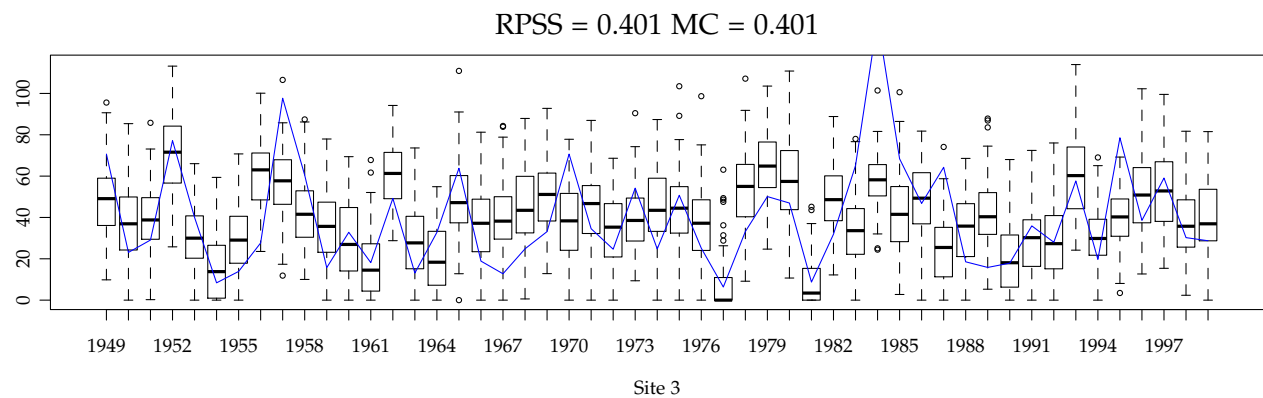
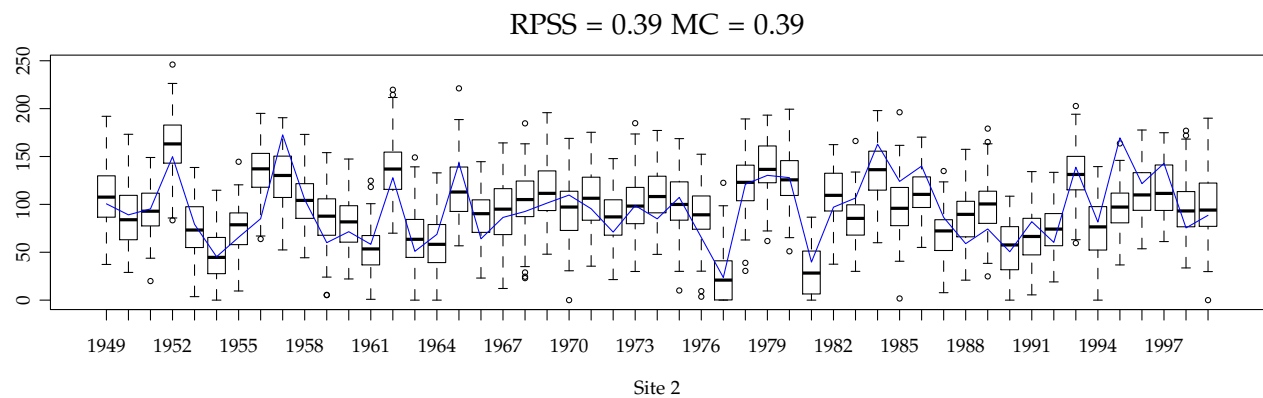
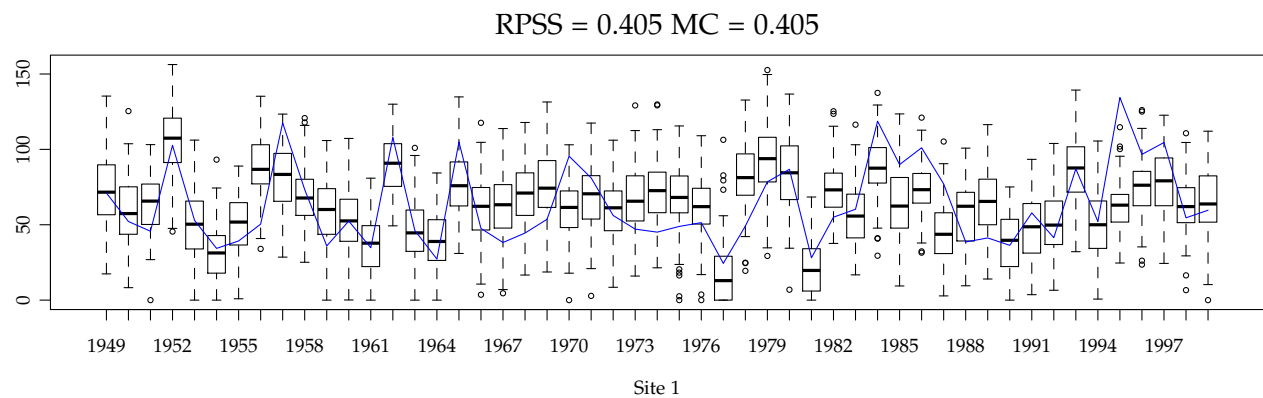


Figure 22: Svd predictions for first three sites

```

> layout(cbind(c(1:3)))
> for(i in 4:6){
+   this.site <- ypred.svd[i,,]
+   this.site <- as.data.frame(this.site)
+   names(this.site) <- 1949:1999
+   boxplot(this.site,xlab=paste('Site',i))
+   lines(flow[,i],col='blue')
+   mtext(paste('RPSS = ',round(stats.svd$rpss[i],3),
+               'MC = ',round(stats.svd$rpss[i],3)),line=1)
+ }

```

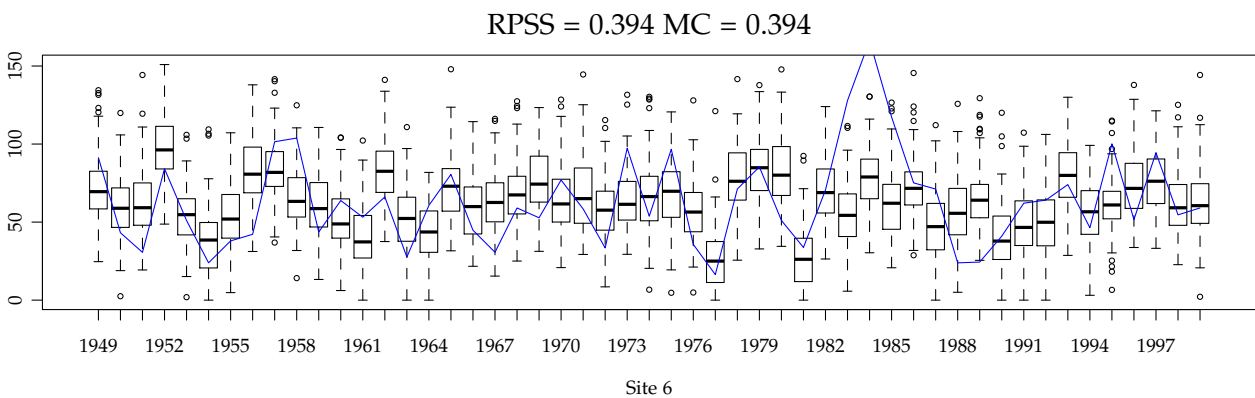
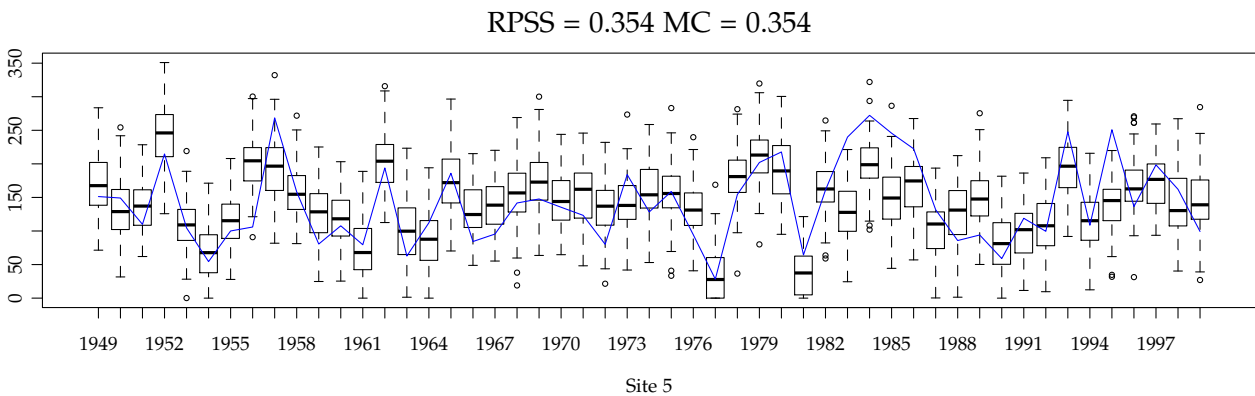
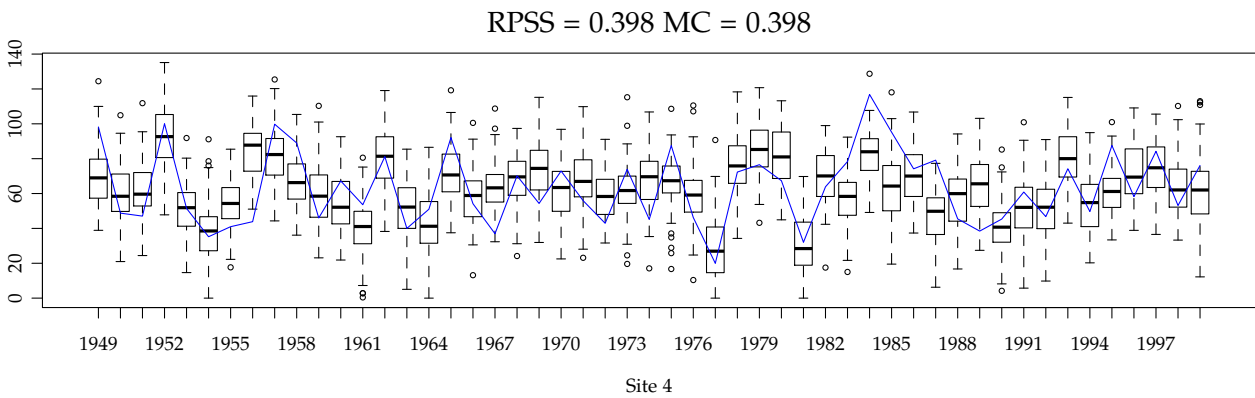


Figure 23: Svd predictions for second three sites

Problem 5

I did not significantly modify the CCA code from the class page so it is not shown here, but it was used to generate forecasts.

```
> layout(cbind(c(1:6)))
> for(i in 1:6){
+   par(mar=c(.9,2,1.5,1))
+   plot(1949:1999, ypred.cca[,i], xlab=paste('Site',i),
+        ylim=range(c(ypred.cca[,i],flow[,i])),main=paste('Site',i),
+        type='l')
+   lines(1949:1999,flow[,i],col='blue')
+ }
```

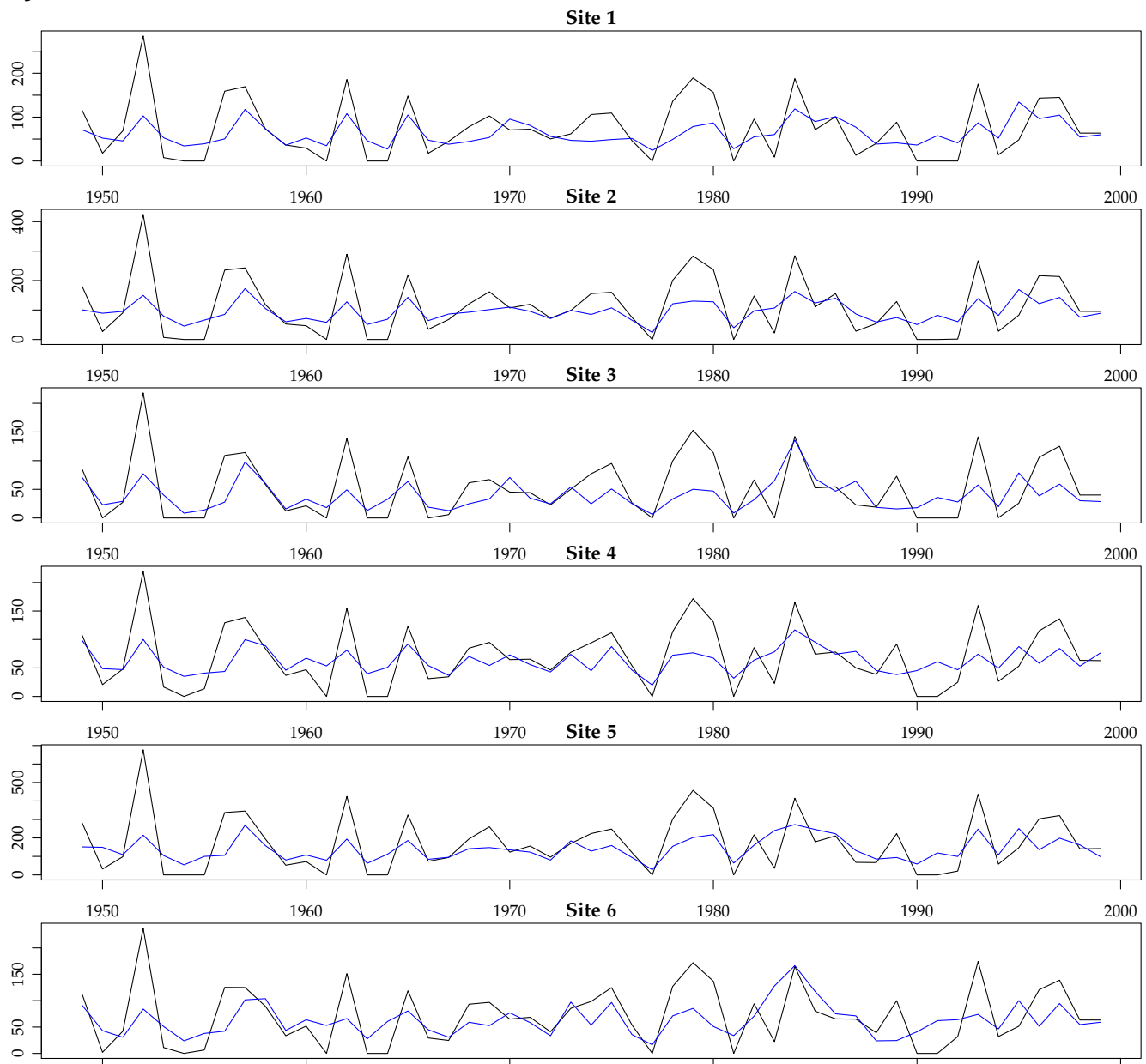


Figure 24: CCA predictions for all sites, historical flow is in blue. Units are thousand cubic meters

Problem 6

Below is the code that does the Markov KNN simulation.

```

source('lib.R')
nyears <- 95
nsim <- nyears * 100
#read data (ac-ft) and convert to cms
x <- as.matrix(read.table('data/Leesferry-mon-data.txt')[,-1])*0.000469050
x.raw <- array(t(x))
x.ann <- x.bin <- ts.annual.mean(ts(x.raw,start=c(1906,1),frequency=12))
n <- length(x.ann)
#historical stats
x.mean <- mean(x.ann); x.sd <- sd(x.ann)
x.skew <- skew(x.ann); x.lag1 <- mylag(x.ann,1,docor=T)

#binary series
x.bin[(x.ann > median(x.ann))] <- 1
x.bin[!(x.ann > median(x.ann))] <- 0

x.wd <- x.ww <- x.dw <- x.dd <- numeric()
p.wd <- p.ww <- p.dw <- p.dd <- n.wd <- n.ww <- n.dw <- n.dd <- 0
for(i in 2:length(x.bin)){
  if(x.bin[i-1] == 0 && x.bin[i] == 0){n.dd <- n.dd + 1; x.dd[n.dd] <- x.ann[i]}
  if(x.bin[i-1] == 0 && x.bin[i] == 1){n.dw <- n.dw + 1; x.dw[n.dw] <- x.ann[i]}
  if(x.bin[i-1] == 1 && x.bin[i] == 0){n.wd <- n.wd + 1; x.wd[n.wd] <- x.ann[i]}
  if(x.bin[i-1] == 1 && x.bin[i] == 1){n.ww <- n.ww + 1; x.ww[n.ww] <- x.ann[i]}
}
p.wd <- n.wd / (n.wd+n.ww); p.ww <- n.ww / (n.wd+n.ww)
p.dw <- n.dw / (n.dw+n.dd); p.dd <- n.dd / (n.dw+n.dd)

#Simulate
x.sim.state <- x.sim <- numeric(nsim)
x.sim.state[1] <- 1
x.sim[1] <- quantile(x.ann,.75)
for(i in 2:nsim){
  r <- runif(1)
  if(x.sim.state[i-1] == 1){
    x.sim.state[i] <- if(r < p.ww) 1 else 0
    pool <- if(r < p.ww) x.ww else x.wd
  }else{
    x.sim.state[i] <- if(r < p.dd) 0 else 1
    pool <- if(r < p.dd) x.dd else x.dw
  }
  x.sim[i] <- sample(pool,1)
}

#calculate stats
x.sim <- matrix(x.sim,ncol=nyears)
x.sim.stats <- annual.stats(x.sim)

save(x.sim.stats, x.sim, x.ann, x.mean, x.sd, x.skew, x.lag1, nsim, nyears,
     file='output/6.Rdata')
```

Figure 25: Markov KNN resampling code.

```

> layout(rbind(1:4))
> boxplot(x.sim.stats[,1],ylim=range(c(x.sim.stats[,1],x.mean)),xlab='Mean')
> points(x.mean,col='red')
> boxplot(x.sim.stats[,2],ylim=range(c(x.sim.stats[,2],x.sd)),xlab='SD')
> points(x.sd,col='red')
> boxplot(x.sim.stats[,3],ylim=range(c(x.sim.stats[,3],x.skew)),xlab='Skew')
> points(x.skew,col='red')
> boxplot(x.sim.stats[,4],ylim=range(c(x.sim.stats[,4],x.lag1)),xlab='Lag1 Cor')
> points(x.lag1,col='red')

```

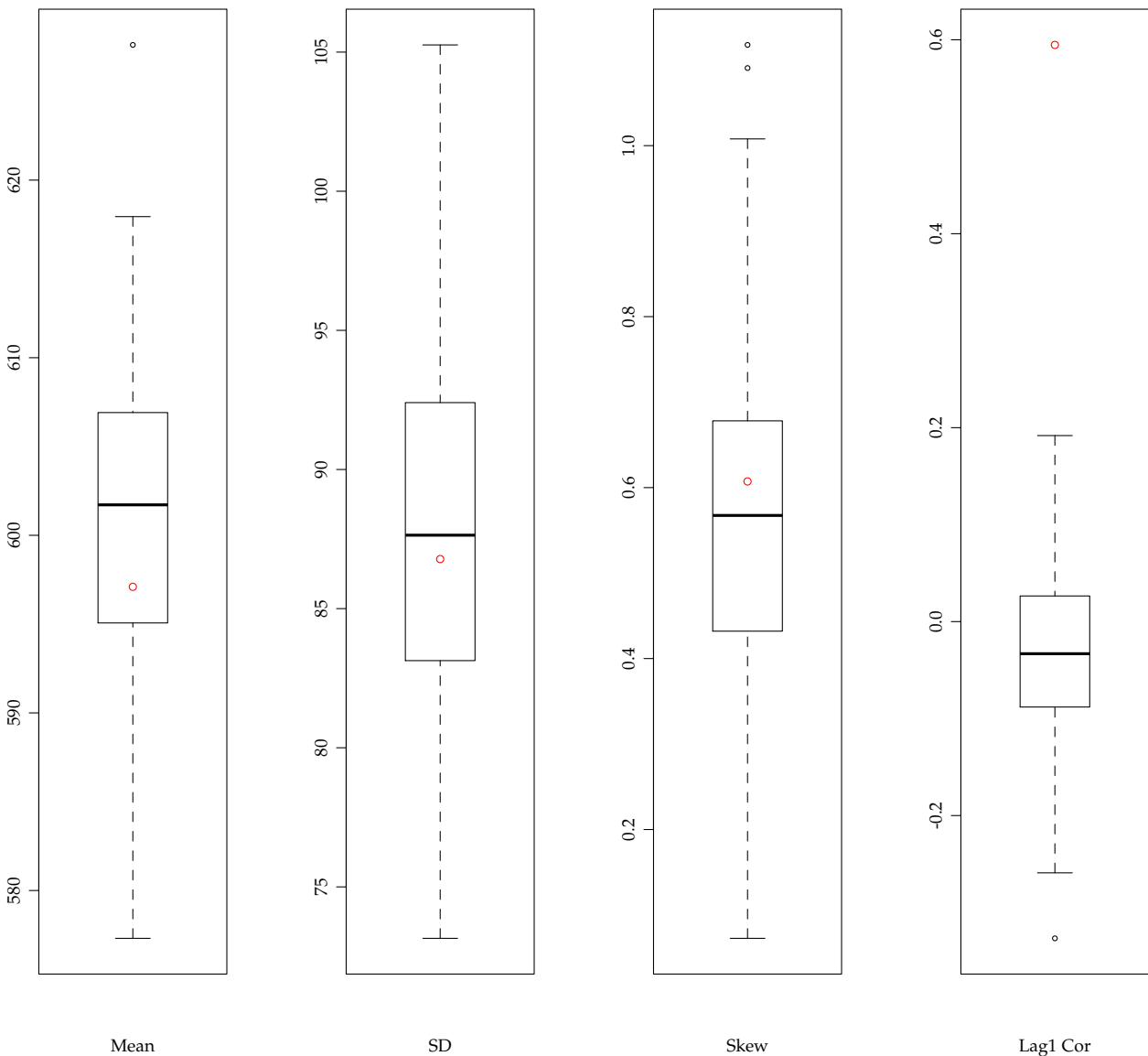


Figure 26: Annual simulated statistics. All statistics are well captured except for Lag 1 autocorrelation. I believe this is due to the way values are sampled from each 'bin.' The only way to get enough sampling variation is to select from all of the values in a bin.

One interesting result I noticed was that the historical pdf was only reproduced when I used a uniform weight function and selected from all the neighbors in a bin. If I used a $1/k$ weight function and set k to \sqrt{n} , the simulated values did not have very much variability at all.

```
> n.pt <- 100
> out <- sm.density(x.ann)
> pt <- seq(min(out$eval.points),max(out$eval.points),length.out=n.pt)
> boxpt <- matrix(NA,nyears,n.pt)
> for(i in 1:nyears){
+   boxpt[i,] <- sm.density(x.sim[i,],
+     eval.points=pt,display='none')$estimate
+   lines(pt,boxpt[i,],col='gray',lwd=.5)
+ }
> sm.density(x.ann,add=T,col='red',lwd=2)
```

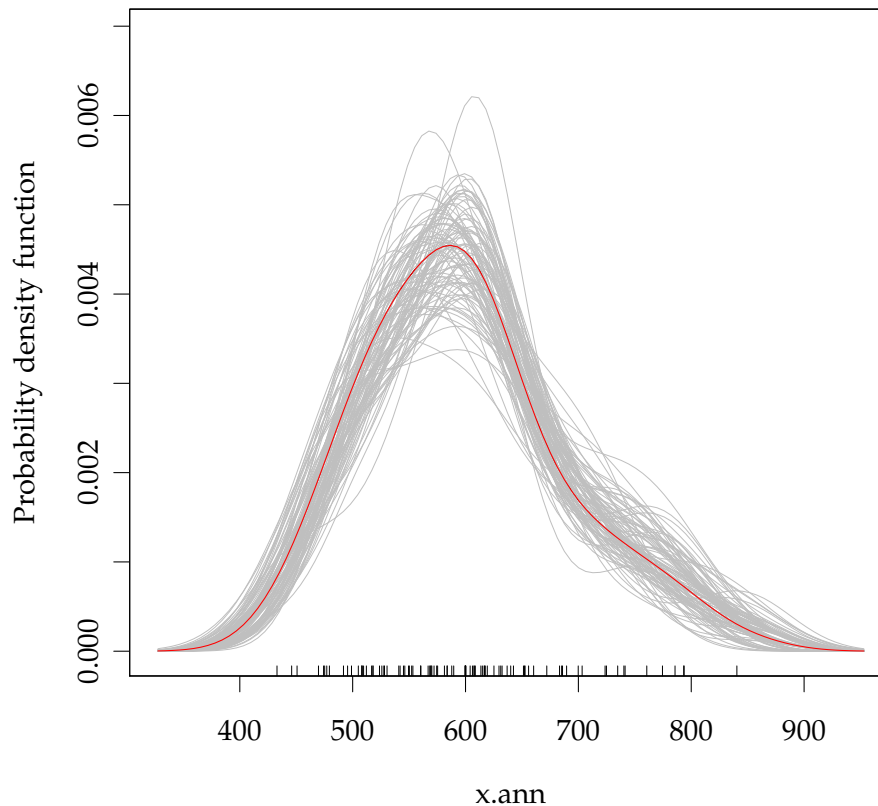


Figure 27: Annual Historical pdf and simulated pdf