









- Or can we just do a model:
 - y=f(lat,lon)+ε
 - y=f(temp(lat,lon),precip(lat,lon))+ε
 - y=f(t)+ε
 - y=f(temp(t))+ε

Space & time different

- Key assumption of GLM is independence of data points
- In space/time there is a particular form of nonindependence:
 - Nearby points are very similar
 - Called autocorrelation (self-correlated)
- World's simplest weather forecast:
 - Tomorrow's weather the same as today
 - Actually has a high degree of accuracy
- Also have a well-defined sense of "distance" between points













In R

co2.diff=diff(co2)
plot(co2.diff)
acf(co2.diff)
acf(co2)
acf(dts)







	Other fancy stuff
	Other failey stuff
N	AO SST <- read.table("Scottish SST.txt", header = T.sep="\t")
s	ST=ts(NAO_SST\$SST,start=1945,freg=12)
Ν	AO=ts(NAO_SST\$NAO,start=1945,freq=12)
С	<pre>cf(stl(NAO,s.window="period") \$time.series[,3], stl(SST,s.window="period" \$time.series[,3])</pre>
d	ucks <- read.table("ducks.txt", header = T, sep = "\t", dec = ".")
1	ibrary(lattice)
x	yplot(Gadwall + Goldeneye + Goosander + Mallard +
	Pochard + Pintail ~ Year, data = ducks,
	type = "l", outer = T, layout = $c(3, 2)$,
	<pre>scales = list(x = list(relation = "same"),</pre>
	<pre>y = list(relation = "free")),</pre>
	<pre>ylab = "Abundance", xlab = "Time (years)")</pre>
1	ibrary(vegan)
d	ucks_pca <- rda(ducks[, -c(1, 2)], scale = T)
р	<pre>lot(ducks_pca, scaling = 2, type = "n", xlab = "axis 1", ylab = "axis 2")</pre>
s	egments(x0 = 0,
	у0 = 0,
	<pre>x1 = scores(ducks_pca, display = "species", scaling = 2)[, 1],</pre>
	<pre>y1 = scores(ducks_pca, display = "species", scaling = 2)[, 2])</pre>
t	ext(ducks_pca, display = "sp", scaling = 2, col = 2)
+	ext(ducks nea display = "wa" labels = ducks[1] scaling = 2 nch = 19







Regression with timeseries

d<-read.table("Hawaii.txt",h=T)
str(d)
d\$N<-sqrt(d\$Moorhen.Kauai)
plot(d\$N~d\$Year)
m<-lm(N-Rainfall+Year,data=d,na.action=na.omit)
plot(m)
plot(residuals(m))
acf(residuals(m))
#uh oh!
m.gls<gls(N-Rainfall+Year,na.action=na.omit,data=d,correlation=corAR1(
form=~Year))
summary(m.gls)
AIC(m)
AIC(m.gls)</pre>

Example

d=read.table('dickabund.csv',h=T, sep=",")
dts=ts(d,start=1966)
acf(dts,lag.max=20)
m=gls(abund~tmin+tmax,data=d,
 corr=corAR1(0.4),method="ML")
m
summary(m)

























library(spatial)
library(MASS)

data(topo)

Polynomial

mpo=surf.ls(degree,data) #deg=2 for quadratic

LOESS

mlo=loess(z~x*y,data=?,degree=2,span=0.25, normalize=F)





















Regression with spatial & exogenous factors

library(AED) data(Boreality) #boreal species over all species Boreality\$Bor<-sqrt(1000*(Boreality\$nBor+1)/Boreality\$nTot) str(Boreality) library(akima) zz<-interp(Boreality\$x,Boreality\$y,Boreality\$Bor)</pre> image(zz) contour(zz,add=T) filled.contour(zz) #simple model m<-lm(Bor~Wet,data=Boreality)</pre> summary(m)library(gstat) E<-rstandard(m) #residuals mydata<-data.frame(E,Boreality\$x,Boreality\$y)</pre> coordinates(mydata)<-c("Boreality.x","Boreality.y")</pre> bubble(mydata,"E",col=c("black","grey")) #errors non-independent!

variogram(Bor~Wet,~x+y,data=Boreality)

In R II

library(spatial) #different form of variogram +correlogram m0<-surf.ls(0,Boreality\$x,Boreality\$y,Boreality\$Bor) variogram(m0,300) correlogram(m0,300) library(nlme) m<-</pre>

gls(Bor~Wet,correlation=corSpher(form=~x+y,nugget=TRUE),data=Bo reality)

#a 3rd form of variogram that works with nlme/gls v<-Variogram(m,form=~x+y,robust=TRUE,maxDist=2000) plot(v)





Point processes

- Record location (x,y coordinates) of "events"
 E.g. plant growing
- Idea of overdispersed/random/clumped
- Poisson process (rate=λ)
 - K(t) = # points within distance t/λ
 - If Poisson K(t)=πt²
 - Study L(t)=sqrt(K(t)/π) should be straight line
- Great deal of machinery
- Also works on quadrats count in each quadrat is Poisson distributed
- Alternatives to Poisson give clumped/overdispersed

library(spatstat)
d<-read.table("ragwortmap2006.txt",h=T)
ragwort< ppp(d\$xcoord,d\$ycoord,c(0,3000),c(0,1500),marks=d\$type)
plot(ragwort)
summary(ragwort)
plot(quadratcount(ragwort))
Z <- density.ppp(ragwort)
plot(Z,main="")
K <- Kest(ragwort)
plot(K) #note observed (black)>red null→clumped
pc <- pcf(ragwort)
plot(pc)</pre>













