

Temporal autocorrelation

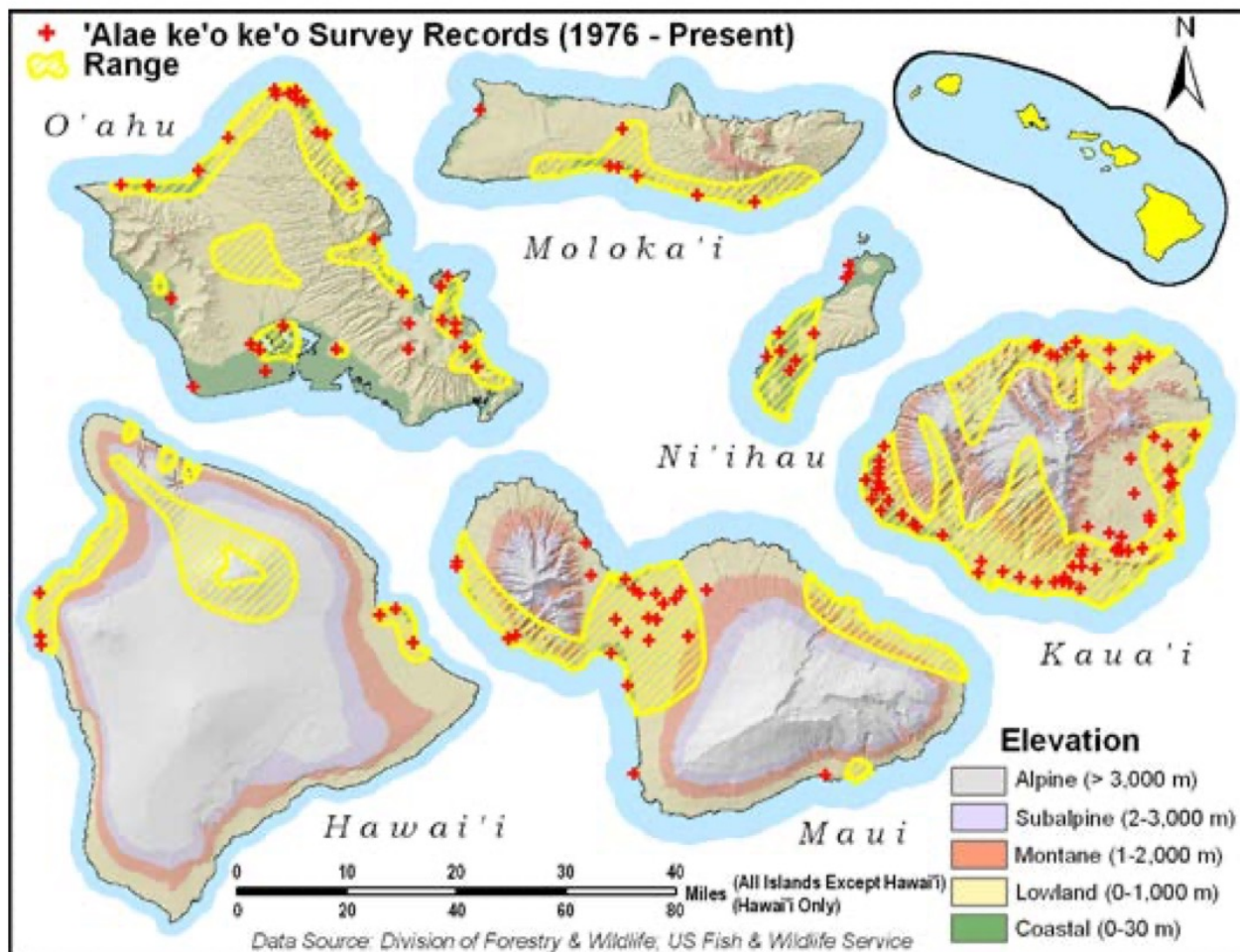
Lots of ways to analyze/model time series data:

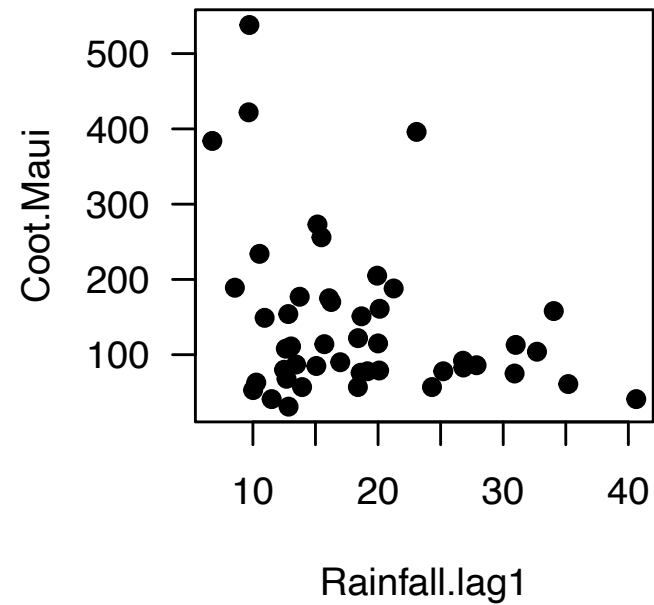
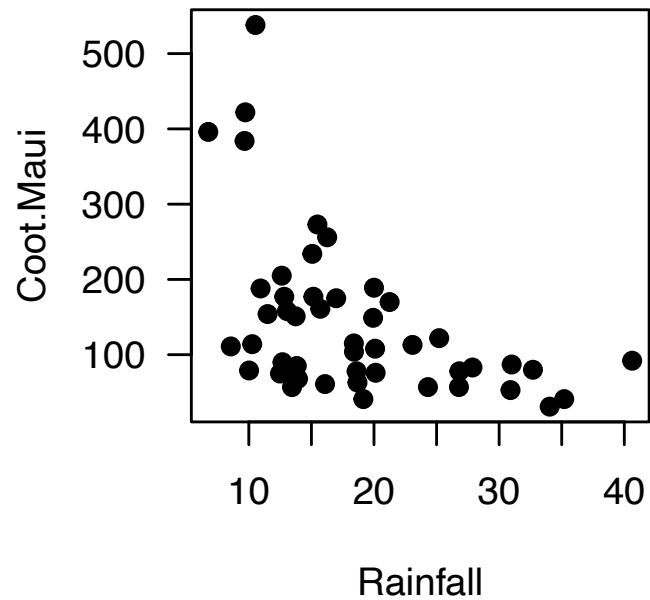
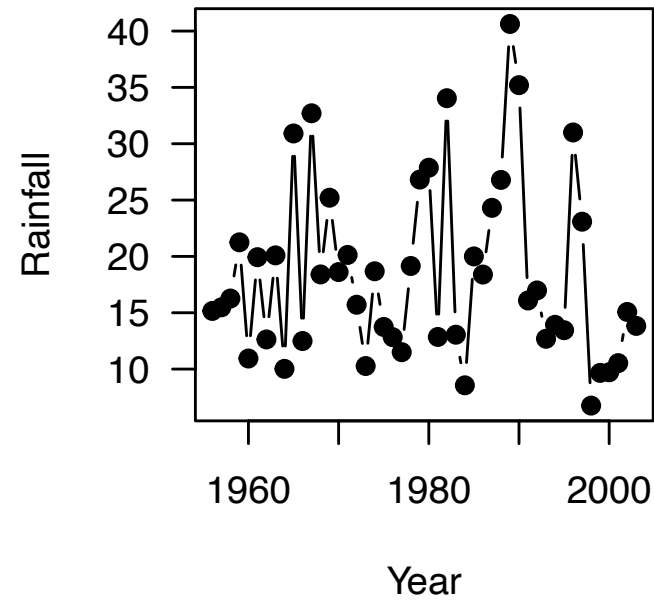
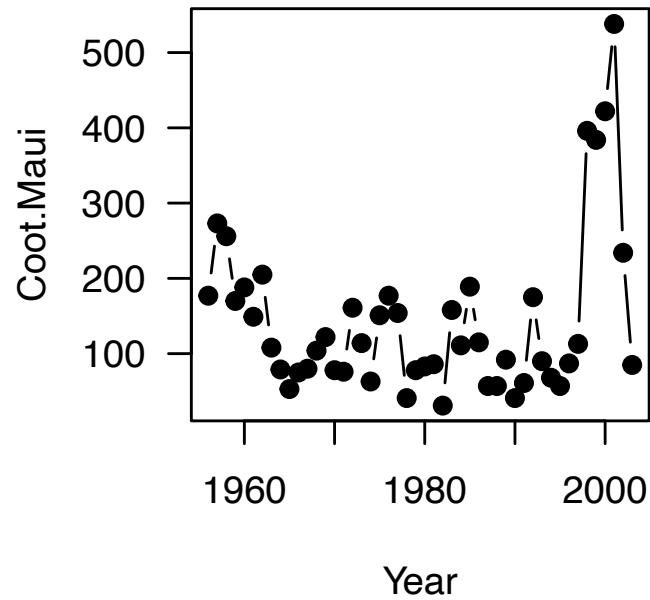
- Autoregressive
- Spectral analysis
- Mechanistic population dynamics

How to use time series data in LM, GLM, GLMM, etc?

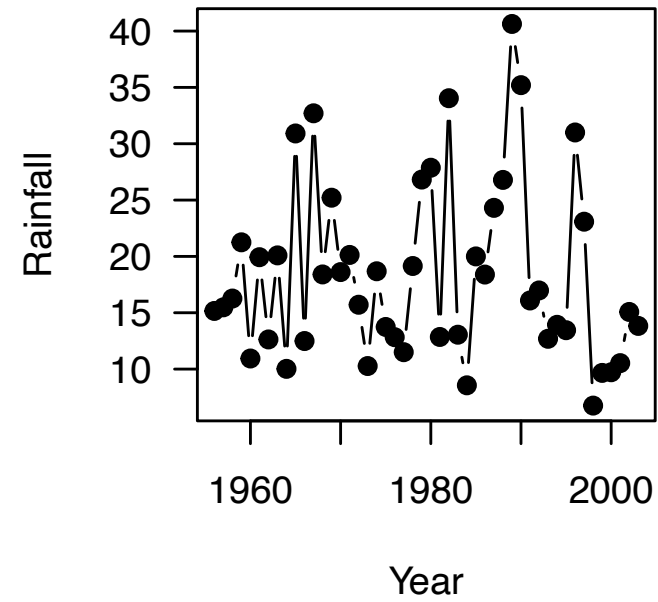
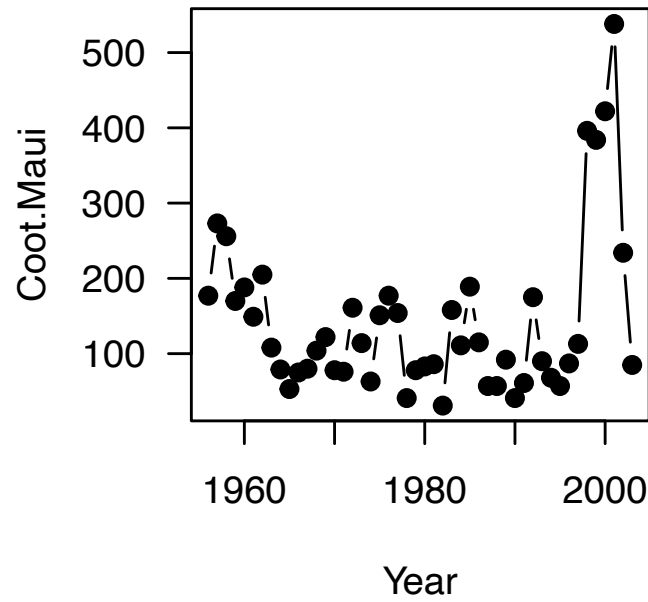
'alae ke'oke'o - Hawaiian coot - *Fulica alai*

Endemic and endangered





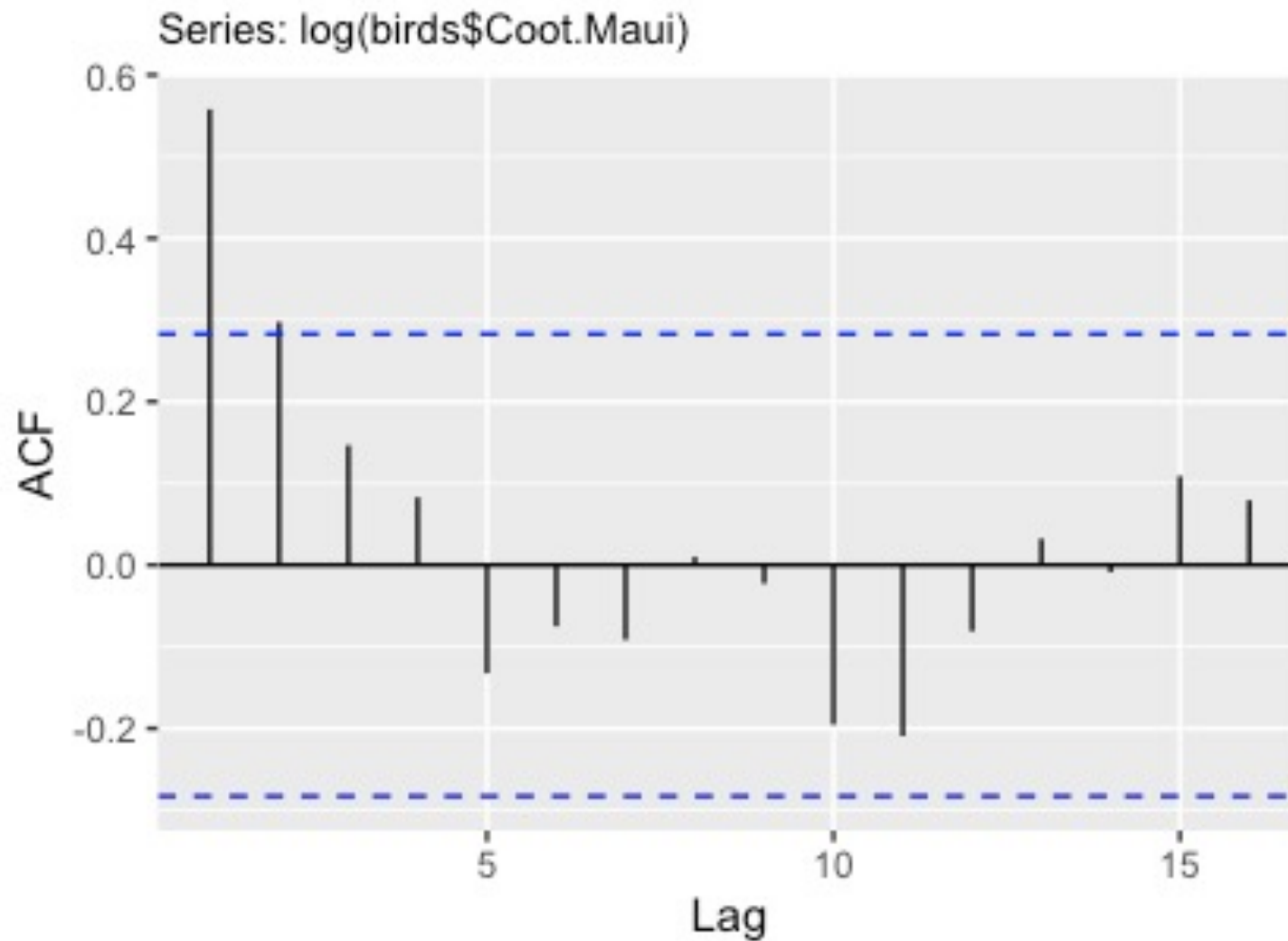
- Rainfall may affect habitat availability, but also counting/detection success
- Treat the bird counts as lognormal



Patterns in time series can be explore with autocorrelation function:

$$r_k = \frac{\sum_{t=1}^{N-k} (Y_t - \bar{Y})(Y_{t+k} - \bar{Y})}{\sum_{t=1}^N (Y_t - \bar{Y})^2}$$

```
ggAcf(log(birds$Coot.Maui))
```



Why might abundance be correlated at 1-2 year lag?

How can we model this kind of dependence, using only the time series itself?

Autoregressive model

The observation at time t depends on one or more previous observations

$$Y_t = c + \varphi Y_{t-1} + \varepsilon_t$$

Usually assume φ is between -1 and 1; represents the lag-1 autocorrelation

This is **AR(1)**

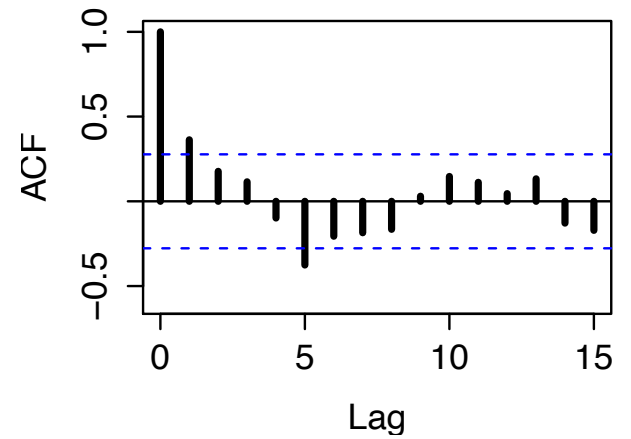
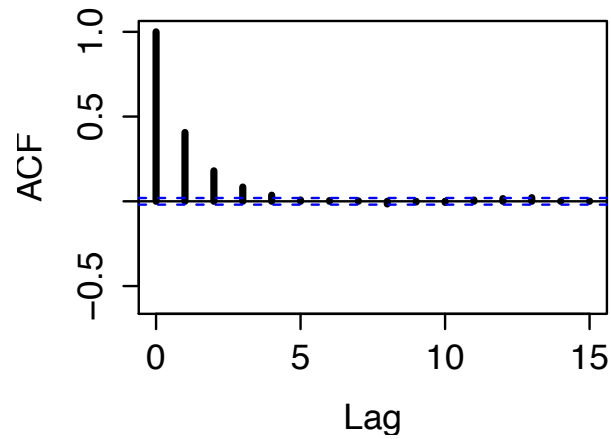
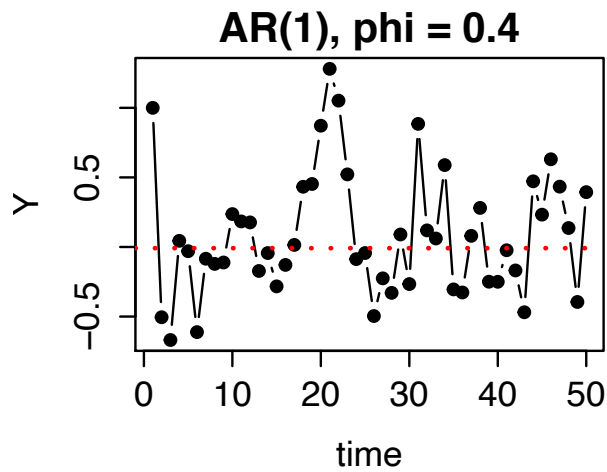
AR(2):

$$Y_t = c + \varphi_1 Y_{t-1} + \varphi_2 Y_{t-2} + \varepsilon_t$$

#AR1

```
phis = c(0.4, 0.8, -0.5)
par(mfrow = c(3,3), mar = c(4,4,2,1))
for (j in 1:3) {
  Y = 1
  tmax = 10000
  SD = 0.5
  phi = phis[j]
  for (t in 2:tmax) {
    Y[t] = phi*Y[t-1] + rnorm(1, mean = 0, sd = SD)
  }

  plot(c(1:50), Y[1:50], type = 'b', pch = 19, main = paste('AR(1), phi =', phi),
       xlab = 'time', ylab = 'Y')
  abline(h = mean(Y), col = 'red', lty = 3, lwd = 2)
  acf(Y, lwd = 3, lag.max = 15, ylim = c(-0.3, 1), main = '10000 observations')
  acf(Y[1:50], lwd = 3, lag.max = 15, ylim = c(-0.3, 1), main = '50 observations')
}
```

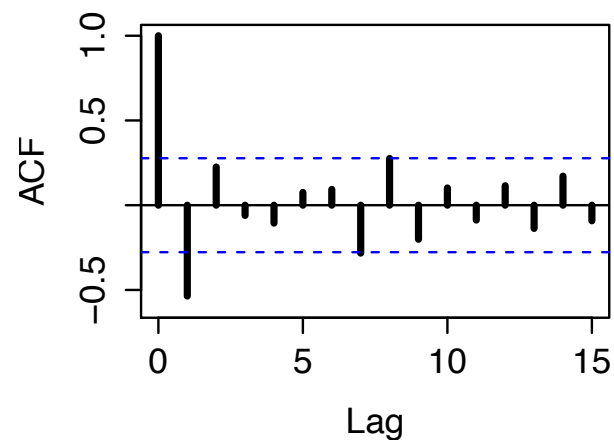
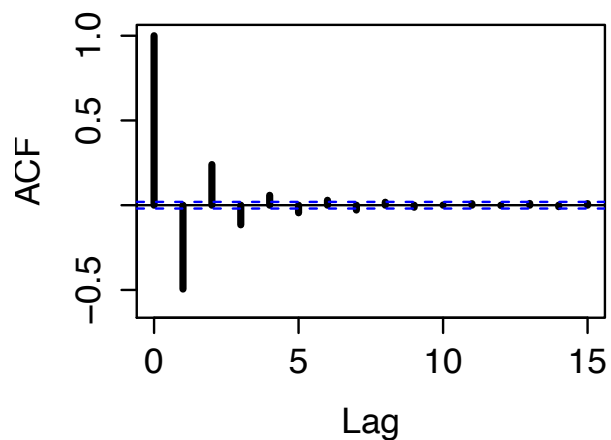
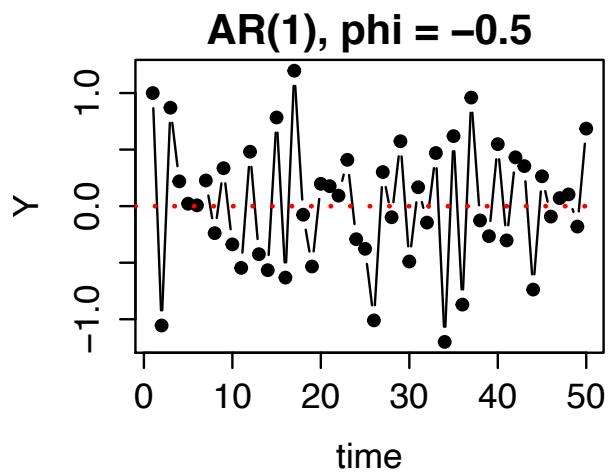
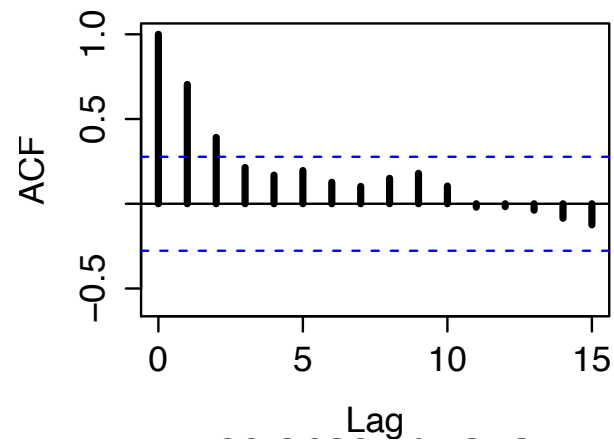
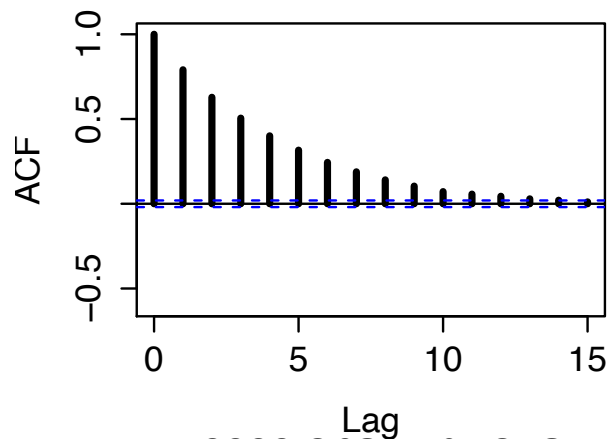
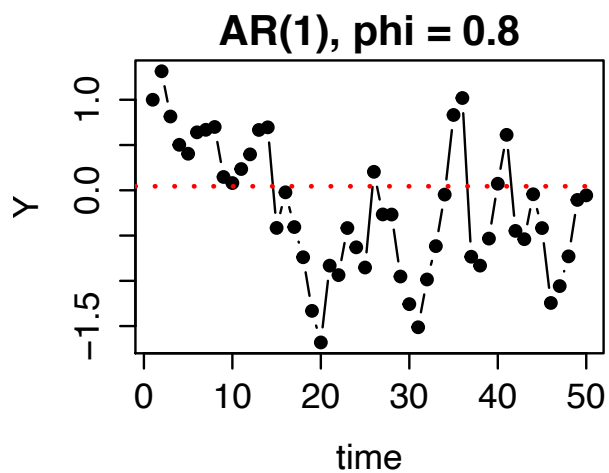
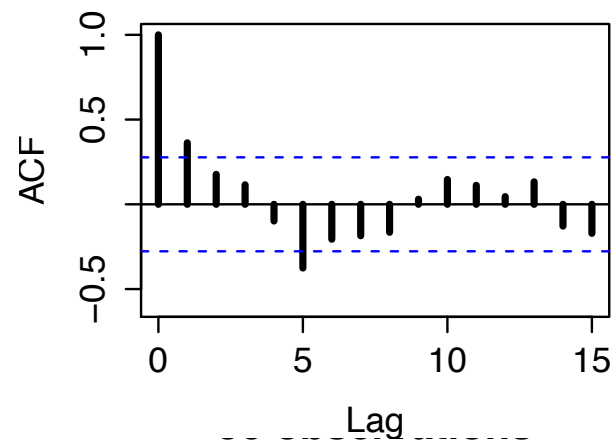
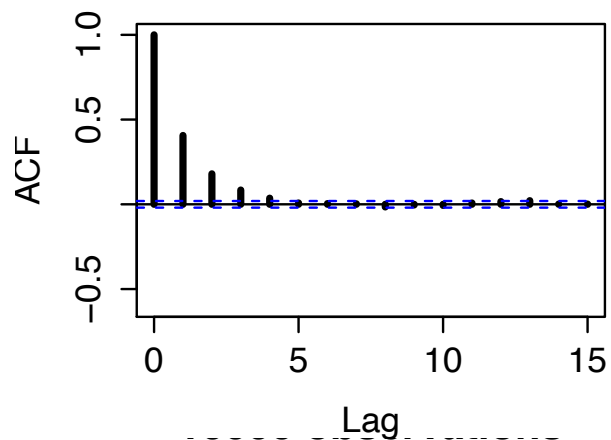
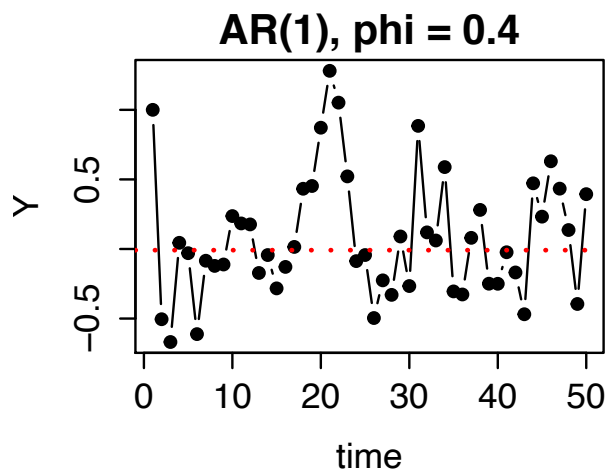


$$Y_2 = \phi Y_1 + \varepsilon_2$$

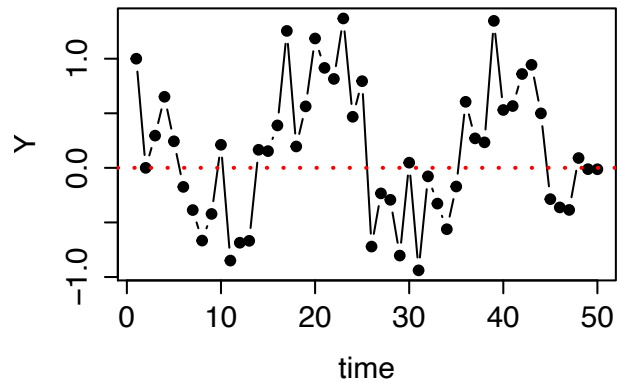
$$Y_3 = \phi Y_2 + \varepsilon_3 = \phi(\phi Y_1 + \varepsilon_2) + \varepsilon_3 = \phi^2 Y_1 + \phi \varepsilon_2 + \varepsilon_3$$

Lag-1	0.4
Lag-2	$0.4^2 = 0.16$
Lag-3	$0.4^3 = 0.064$

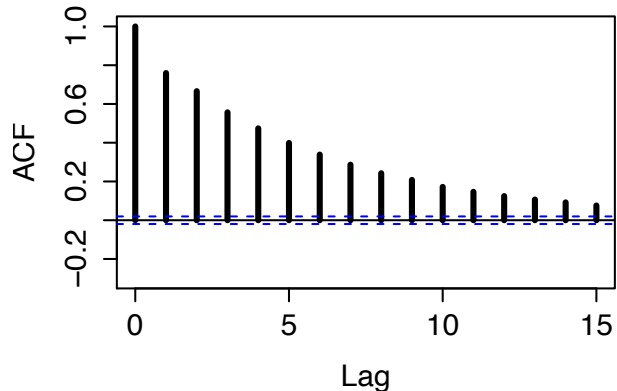
E.g., a species that lives for a few years



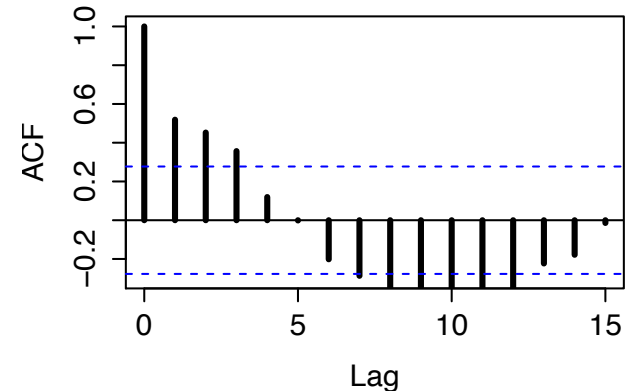
AR(2), $\phi_1 = 0.6$, $\phi_2 = 0.2$



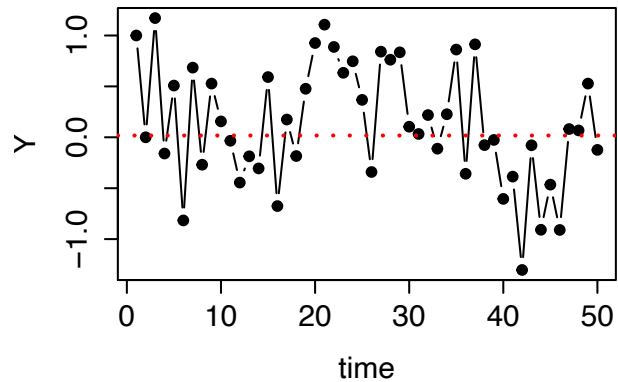
10000 observations



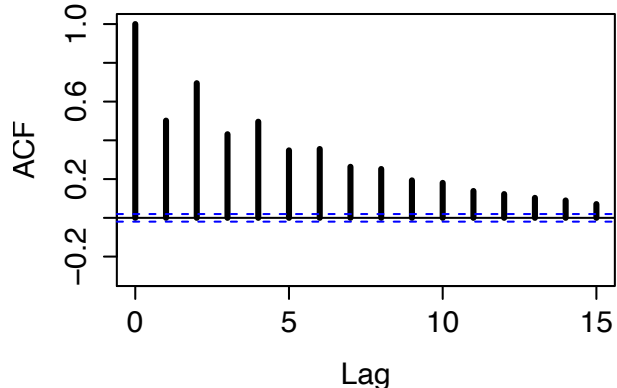
50 observations



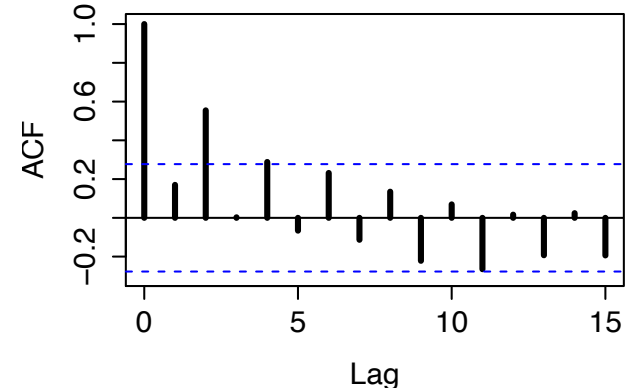
AR(2), $\phi_1 = 0.2$, $\phi_2 = 0.6$



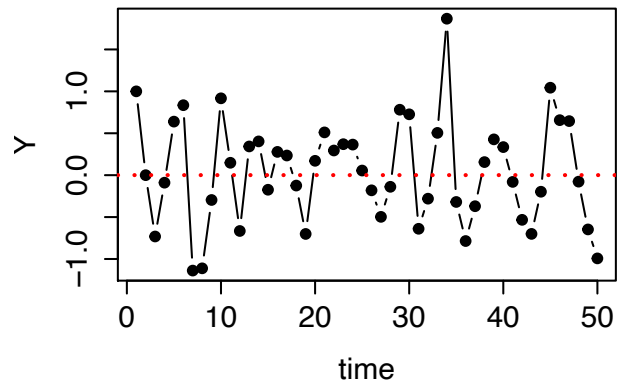
10000 observations



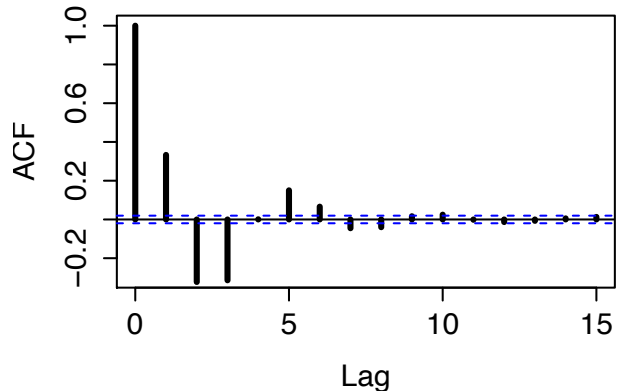
50 observations



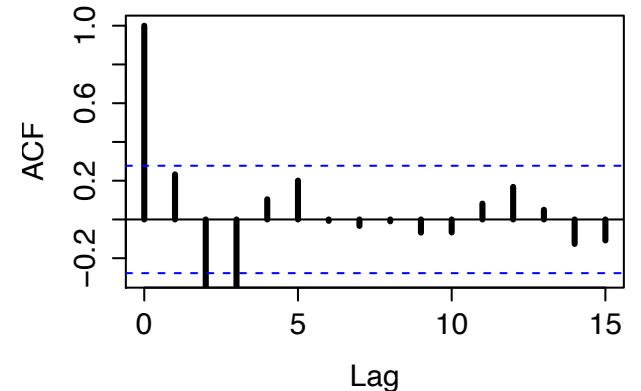
AR(2), $\phi_1 = 0.5$, $\phi_2 = -0.5$



10000 observations



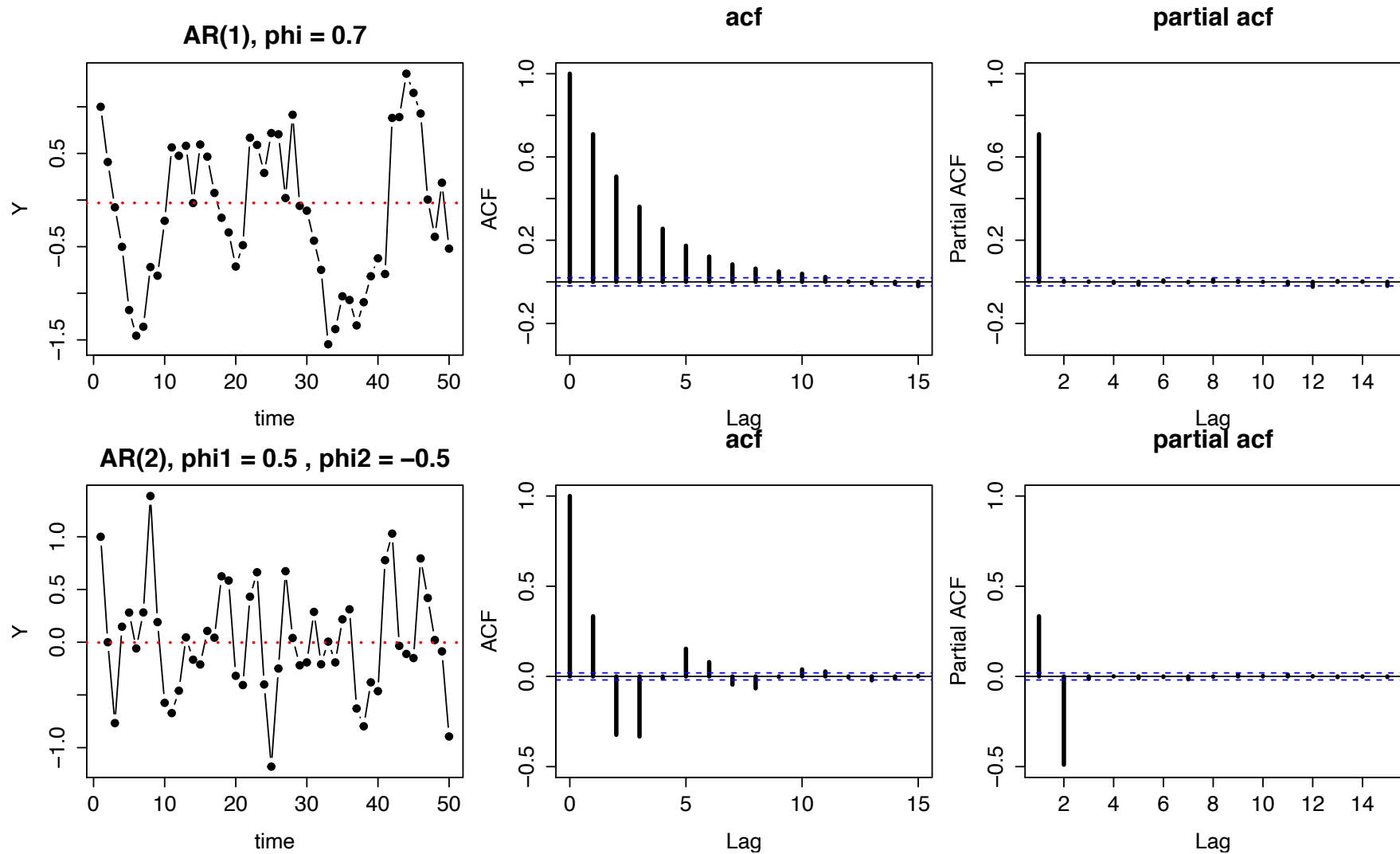
50 observations



Partial autocorrelation function:

Correlation at lag k , after removing the effect of earlier lags

`acf(mydata, type = "partial")`



Moving average model

Value at time t doesn't depend directly on previous values

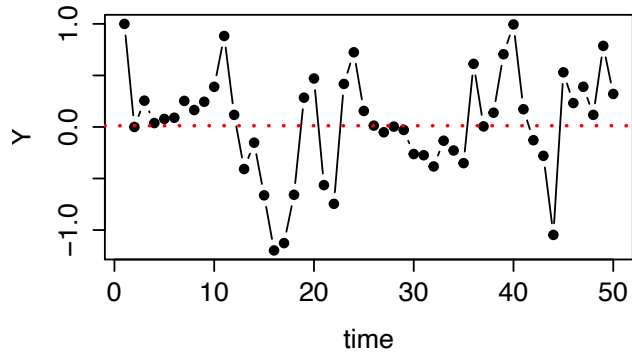
Depends on random 'shocks' from previous timesteps

$$Y_t = \mu + \varepsilon_t + \theta_1 \varepsilon_{t-1} + \cdots + \theta_q \varepsilon_{t-q}$$

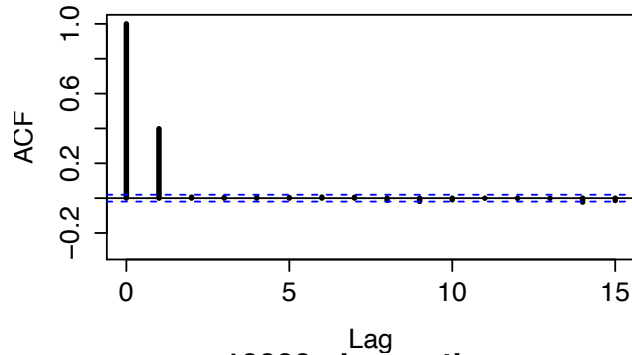
Connection to population dynamics less obvious, but good at modeling short-term autocorrelation that does not have a 'memory'

$$Y_t = \mu + \varepsilon_t + \theta_1 \varepsilon_{t-1} + \cdots + \theta_q \varepsilon_{t-q}$$

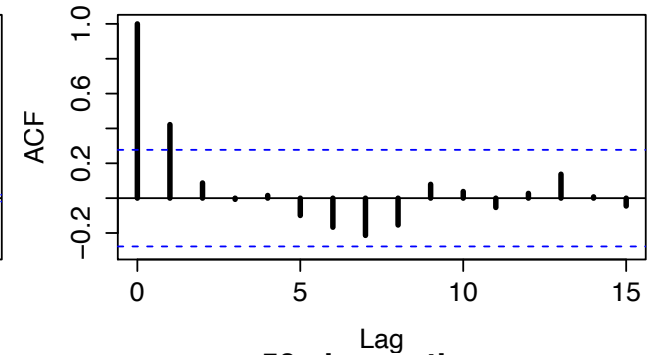
MA(2), theta1 = 0.5 , theta2 = 0



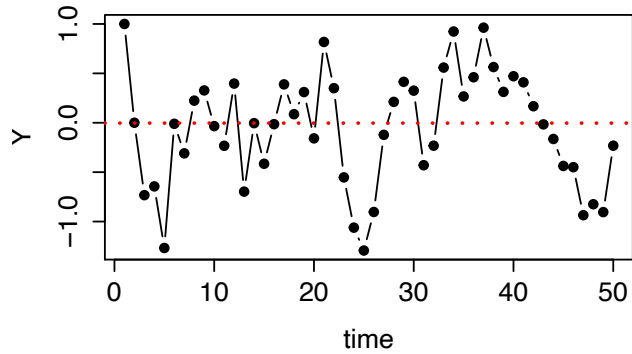
10000 observations



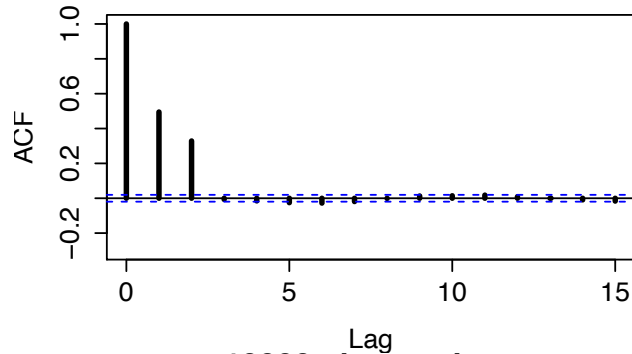
50 observations



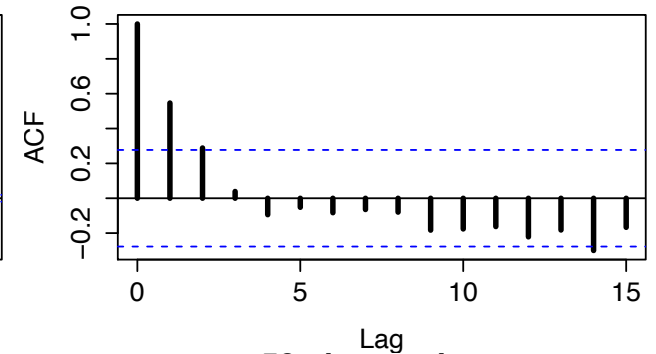
MA(2), theta1 = 0.5 , theta2 = 0.5



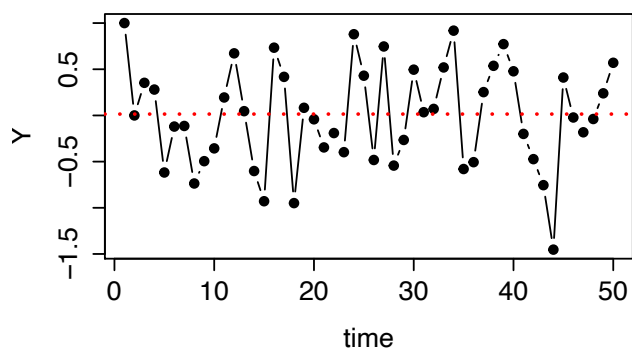
10000 observations



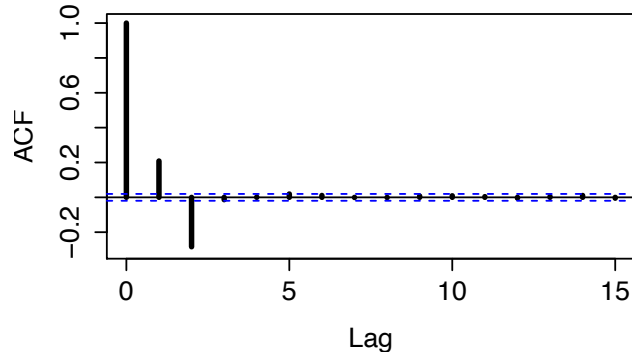
50 observations



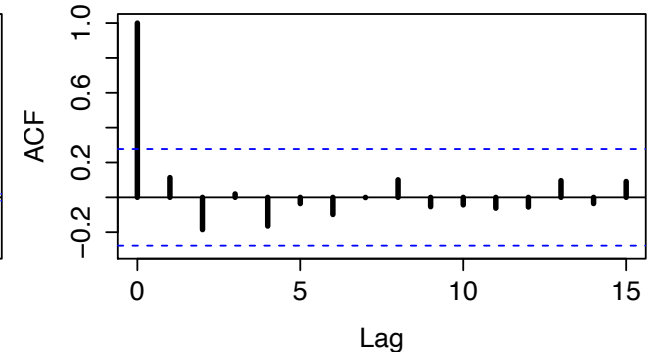
MA(2), theta1 = 0.7 , theta2 = -0.5



10000 observations



50 observations



In principle, can distinguish AR and MA, and their order, using acf

But with limited data it's not so clear. Try simple models, compare, add complexity.

Generalized least squares (GLS)

We want to 1) model the effect of rainfall on abundance, 2) while also accounting for temporal autocorrelation

We will do this by adding a **model for the residuals**

Terminology is unfortunate:

- GLS is for normal data with correlated residuals
- GLM for non-normal data with uncorrelated residuals
- GEE = Generalized Estimating Equations is for non-normal data with correlated residuals

$$Y_t = \beta_0 + \beta_1 X_t + \varepsilon_t$$

$$\begin{bmatrix} \varepsilon_1 \\ \varepsilon_2 \\ \varepsilon_3 \\ \varepsilon_4 \end{bmatrix} = \varepsilon \sim \text{MultivariateNormal}(\Sigma)$$

$$\Sigma = \sigma^2 \begin{bmatrix} 1 & \varphi & \varphi^2 & \varphi^3 \\ \varphi & 1 & \varphi & \varphi^2 \\ \varphi^2 & \varphi & 1 & \varphi \\ \varphi^3 & \varphi^2 & \varphi & 1 \end{bmatrix}$$

Sigma² is the residual variance

Phi is the correlation between consecutive residuals

This is an **AR(1)** model

Note: we assume the amount of variability does not change over time

Stationary: the distribution from which the data are drawn does not change over time

```
mod.nopredictors = gls(log(Coot.Maui) ~ 1, data = birds, correlation =  
corAR1(form =~ Year))
```

Syntax like lm(), but with a correlation argument

See ?corStruct

AR1 has its own function

Can also do AR1 within groups: form =~ Year | Site


```

## Generalized least squares fit by REML
##   Model: log(Coot.Maui) ~ 1
##   Data: birds
##   AIC   BIC logLik
##   82.3 87.85 -38.15
##
## Correlation Structure: AR(1)
## Formula: ~Year
## Parameter estimate(s):
##   Phi
## 0.587
##
## Coefficients:
##               Value Std.Error t-value p-value
## (Intercept) 4.752      0.18    26.4      0
##
## Standardized residuals:
##      Min      Q1      Med      Q3      Max
## -2.01305 -0.61361 -0.08572  0.63491  2.34595
##
## Residual standard error: 0.6547
## Degrees of freedom: 48 total; 47 residual

```

Can compare raw residuals and ‘normalized’ residuals

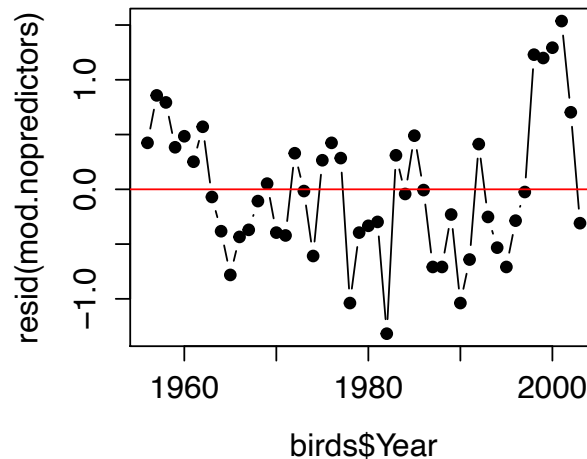
$$Y_t - \text{fitted}(Y_t) = \eta_t = \varphi\eta_{t-1} + \varepsilon_t$$

```

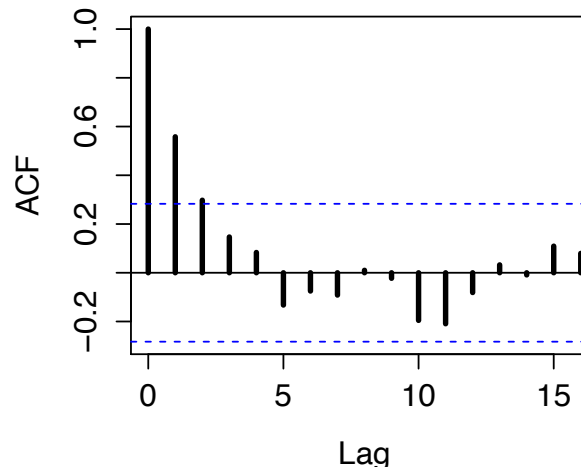
plot(resid(mod.nopredictors) ~ birds$Year, type = 'b', pch = 19, main = 'raw o
ver time')
abline(h = 0, col = 'red')
plot(resid(mod.nopredictors, type = "normalized") ~ birds$Year, type = 'b', pc
h = 19, main = 'normalized over time')
abline(h = 0, col = 'red')
acf(resid(mod.nopredictors), lwd = 3, main = 'raw residuals')
acf(resid(mod.nopredictors, type = "normalized"), lwd = 3, main = 'normalized
residuals')

```

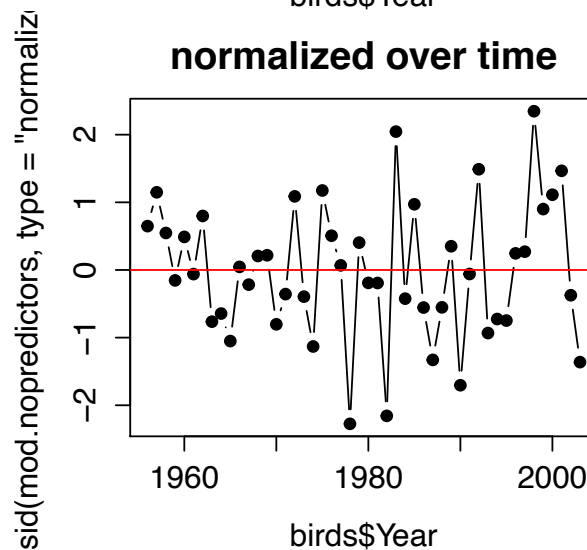
raw over time



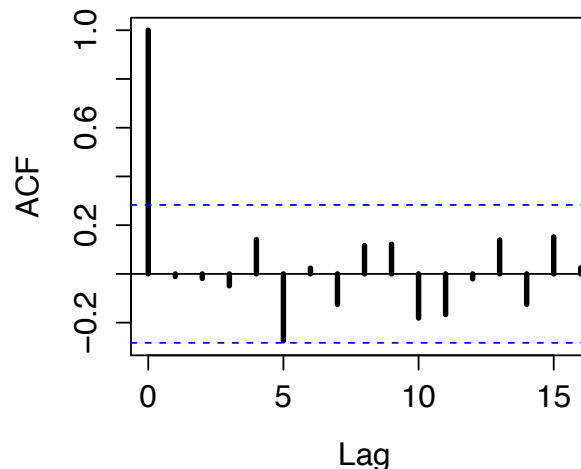
raw residuals



normalized over time



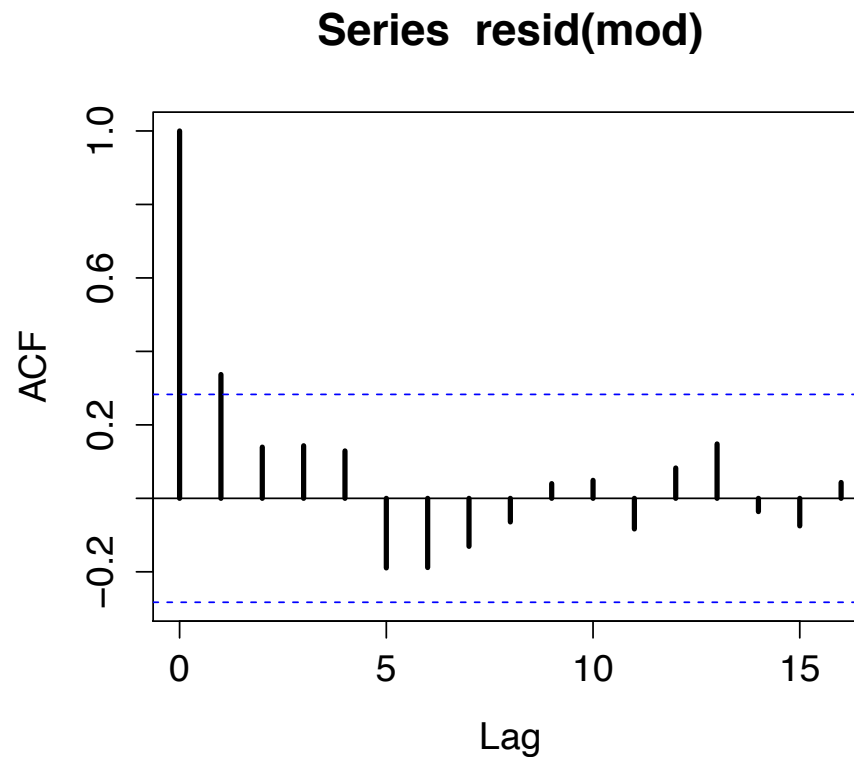
normalized residuals



Important: model assumes residuals independent after accounting for the predictors

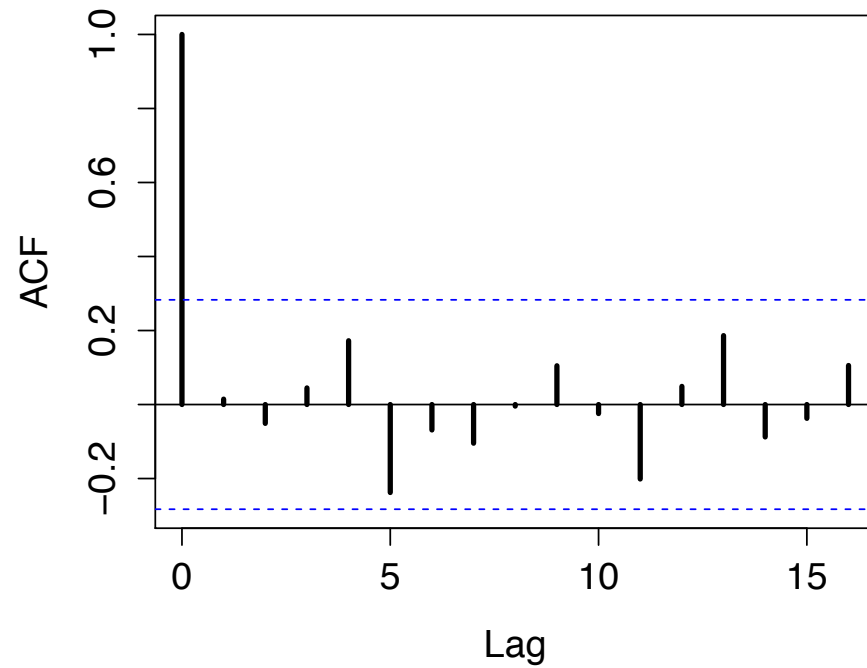
So using the right predictor may be sufficient (like we saw with the spatial smoother)≈

```
mod = lm(log(Coot.Maui) ~ Rainfall, data = birds)
acf(resid(mod))
```



```
mod1 = gls(log(Coot.Maui) ~ Rainfall, data = birds, correlation = corAR1(form  
=~ Year))  
acf(resid(mod1, type = "normalized"))
```

Series resid(mod1, type = "normalized")



```

Model: log(Coot.Maui) ~ Rainfall + Rainfall.lag1
Data: subset(birds, !is.na(Rainfall.lag1))
      AIC      BIC    logLik
74.40192 83.65266 -32.20096

```

Correlation Structure: AR(1)

Formula: ~Year

Parameter estimate(s):

```

      Phi
0.4161663

```

Coefficients:

	Value	Std.Error	t-value	p-value
(Intercept)	5.538649	0.30630520	18.082126	0.0000
Rainfall	-0.032254	0.00949971	-3.395276	0.0015
Rainfall.lag1	-0.010790	0.00950933	-1.134696	0.2626

Correlation:

	(Intr)	Ranfl1
Rainfall	-0.737	
Rainfall.lag1	-0.739	0.294

Standardized residuals:

	Min	Q1	Med	Q3	Max
	-2.05602659	-0.81469555	0.06235979	0.62591168	2.26555967

Residual standard error: 0.5269013

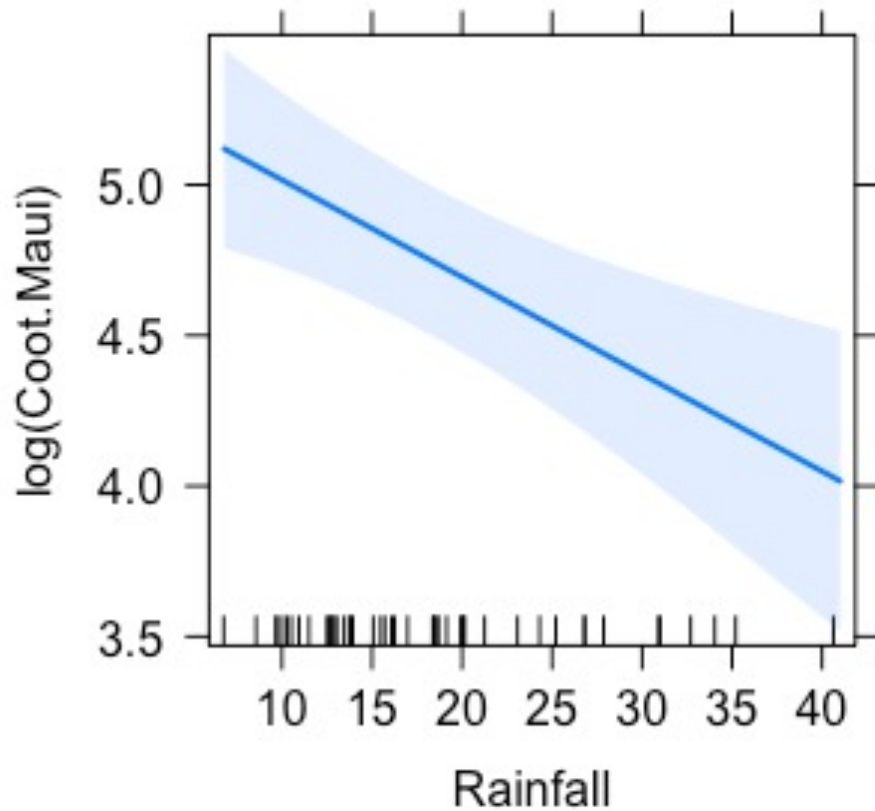
Degrees of freedom: 47 total; 44 residual

```
> anova(mod1)
```

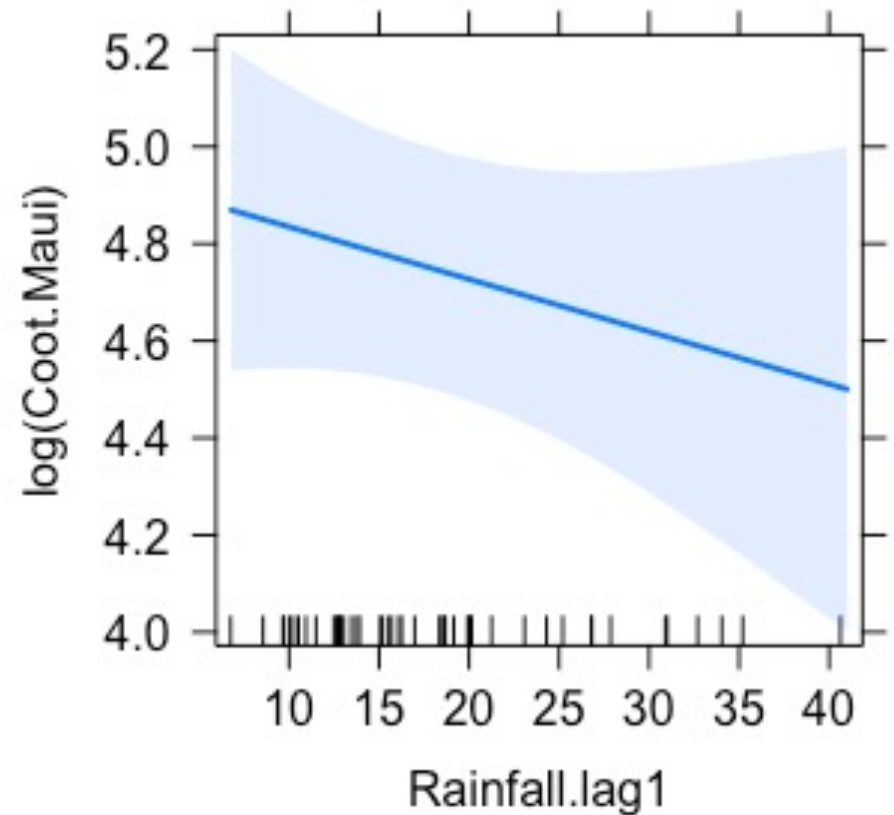
```
Denom. DF: 44
```

	numDF	F-value	p-value
(Intercept)	1	1517.5655	<.0001
Rainfall	1	10.2611	0.0025
Rainfall.lag1	1	1.2875	0.2626

Rainfall effect plot



Rainfall.lag1 effect plot



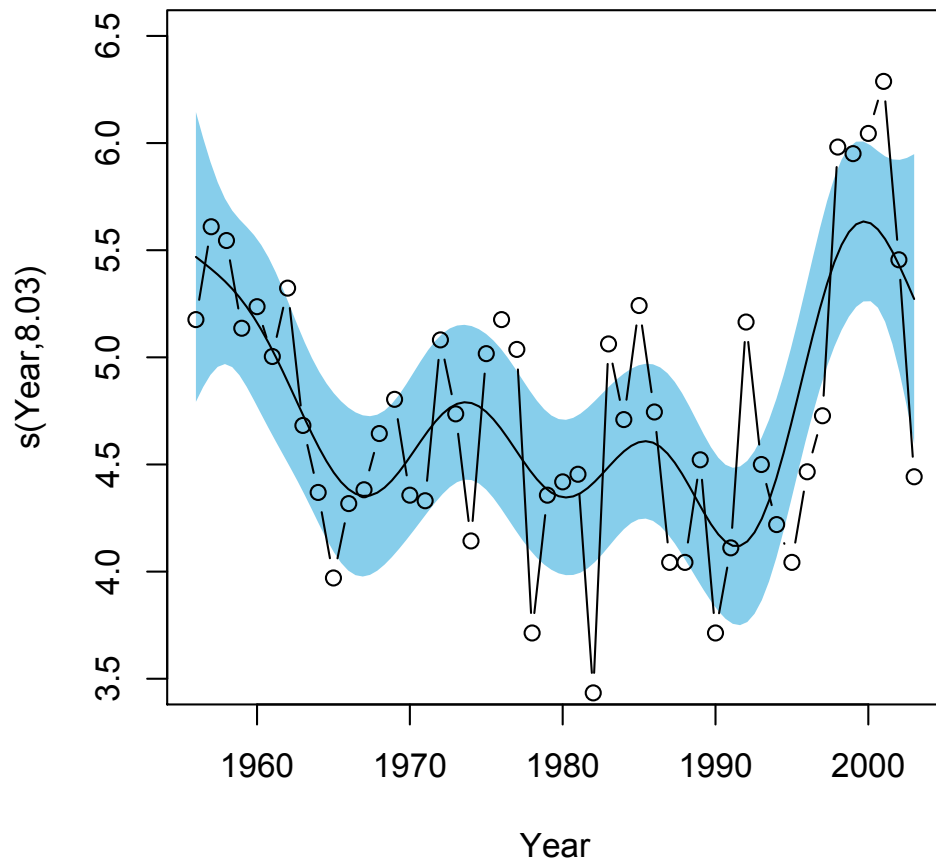
$\exp(5) = 148$ to $\exp(4) = 55$

Conclusions?

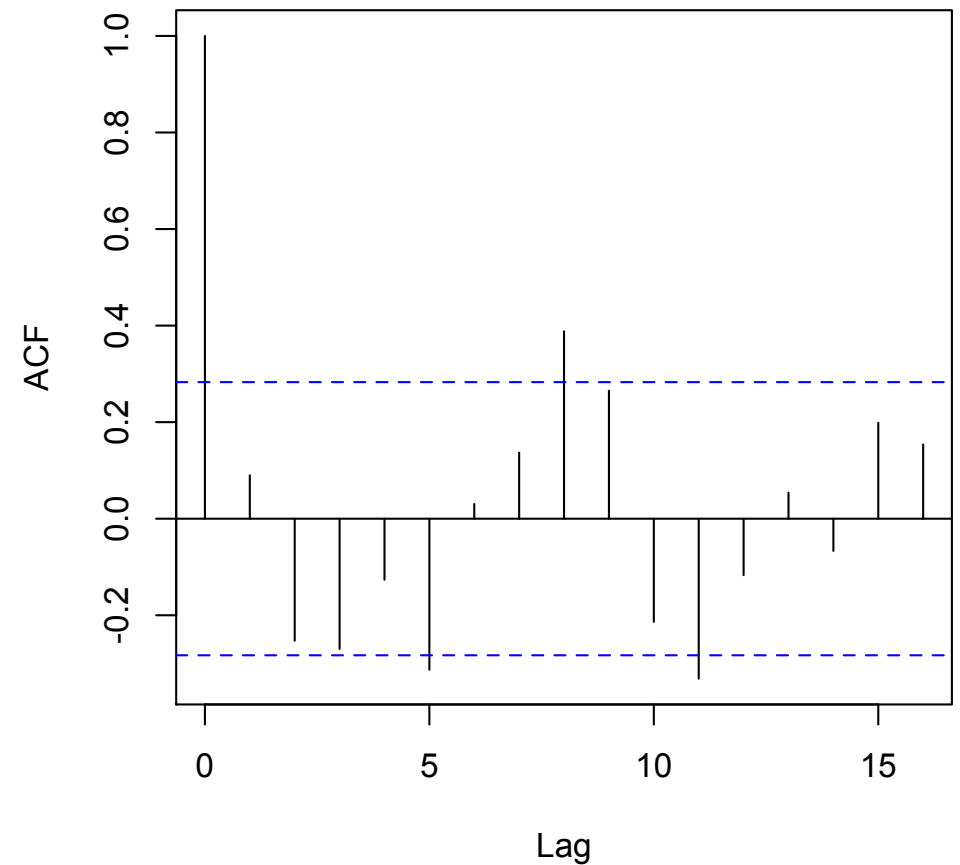
- High rainfall in current year may make detection harder (more habitat)
- High rainfall in previous year doesn't seem to translate into a larger population this year

Alternative to GLS: can we account for temporal autocorrelation using a smoother?

```
mod.gam.year = gam(log(Coot.Maui) ~ s(Year), data = birds)
```



Series resid(mod.gam.year)



Alternative to GLS: can we account for temporal autocorrelation using a smoother?

```
mod.gam.year = gam(log(Coot.Maui) ~ s(Year), data = birds)
```

```
> gam.check(mod.gam.year)
```

Method: GCV Optimizer: magic

Smoothing parameter selection converged after 9 iterations.

The RMS GCV score gradient at convergence was 4.240182e-07 .

The Hessian was positive definite.

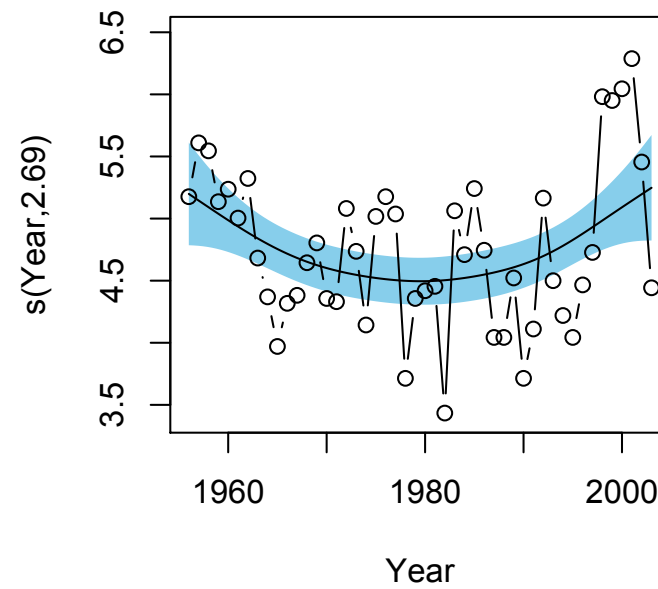
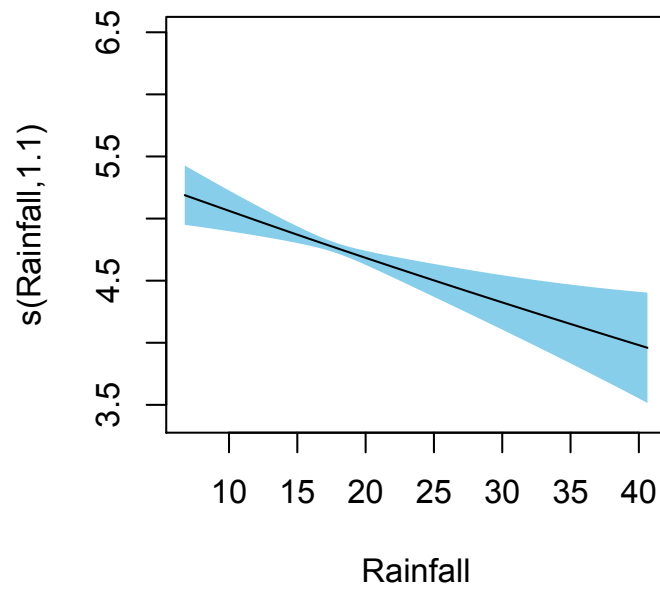
The estimated model rank was 10 (maximum possible: 10)

Model rank = 10 / 10

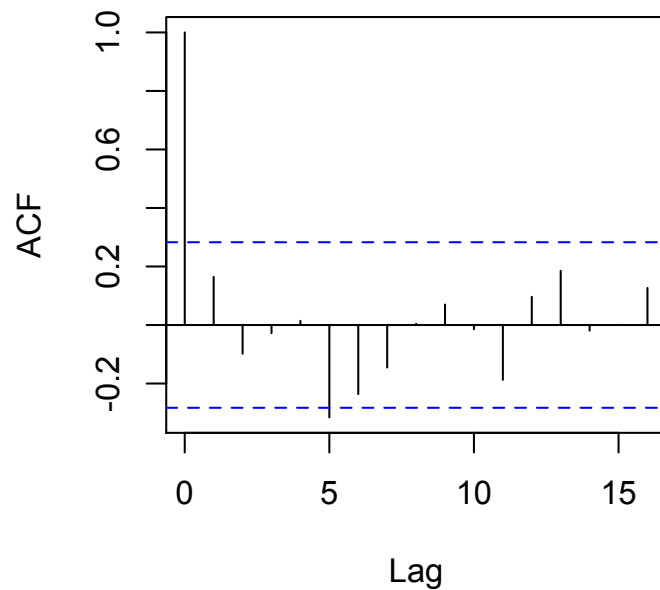
Basis dimension (k) checking results. Low p-value (k-index<1) may indicate that k is too low, especially if edf is close to k'.

	k'	edf	k-index	p-value
s(Year)	9.000	8.033	0.886	0.14

Alternative to GLS: can we account for temporal autocorrelation using a smoother?



Series resid(mod.gam.rainfall)



Alternative to GLS: can we account for temporal autocorrelation using a smoother?

```
mod.gam.rainfall = gam(log(Coot.Maui) ~ s(Rainfall) + s(Year), data = birds)
```

Family: gaussian

Link function: identity

Formula:

```
log(Coot.Maui) ~ s(Rainfall) + s(Year)
```

Parametric coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	4.7485	0.0694	68.43	<2e-16 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	F	p-value
s(Rainfall)	1.102	1.197	12.200	0.000559 ***
s(Year)	2.686	3.349	3.544	0.018407 *

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.437 Deviance explained = 48.2%

GCV = 0.25677 Scale est. = 0.23116 n = 48

For non-normal data: Generalized Estimating Equations

Basically an extension of quasipoisson, quasibinomial

For Poisson, assume the variance is proportional to the mean, modified by the residual correlation structure you specify

```
library(geepack)

birds$idvar = factor(rep('1', nrow(birds)))

gee.rainfall = geeglm(Coot.Maui ~ Rainfall, family = poisson, id = idvar, corstr = 'ar1',
data = birds)
```

For non-normal data: Generalized Estimating Equations

Basically an extension of quasipoisson, quasibinomial

For Poisson, assume the variance is proportional to the mean, modified by the residual correlation structure you specify

Call:

```
geeglm(formula = Coot.Maui ~ Rainfall, family = poisson, data = birds,  
        id = idvar, corstr = "ar1")
```

Coefficients:

	Estimate	Std.err	Wald	Pr(> W)	
(Intercept)	5.45e+00	3.95e-07	1.90e+14	<2e-16	***
Rainfall	-2.95e-02	1.77e-08	2.78e+12	<2e-16	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Estimated Scale Parameters:

	Estimate	Std.err
(Intercept)	54.9	1.27e-06

Correlation: Structure = ar1 Link = identity

Estimated Correlation Parameters:

	Estimate	Std.err
alpha	0.553	1.55e-07

Number of clusters: 1 Maximum cluster size: 48