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Generalized additive models

GAM

GAMs

Statistical power (prediction) vs. ecological interpretation



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Generalized linear and generalized additive models in studies of species distributions: setting the scene

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Generalized additive models

- Linear model (LM)
 - Models response variable using explanatory variables that are associated to the response variable using **specific parametric functions**
 - Linear, log, quadratic
- Generalized additive models (GAM)
 - Models response variable using explanatory variables that are fitted as arbitrary smoothed functions using **non-parametric smoothers**

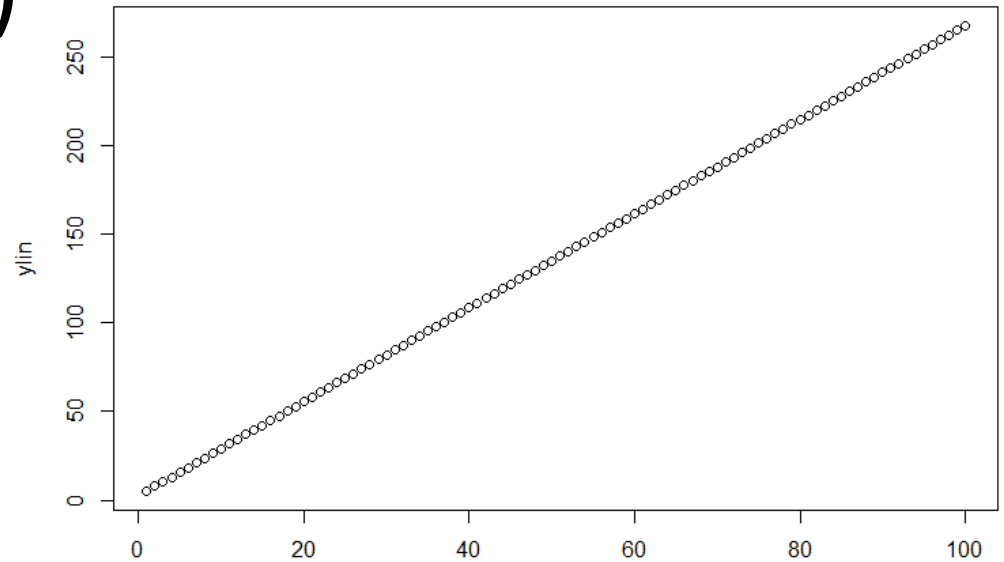
Linear models (lm)

linear parametric functions

```
x <- seq(1,100,1)
```

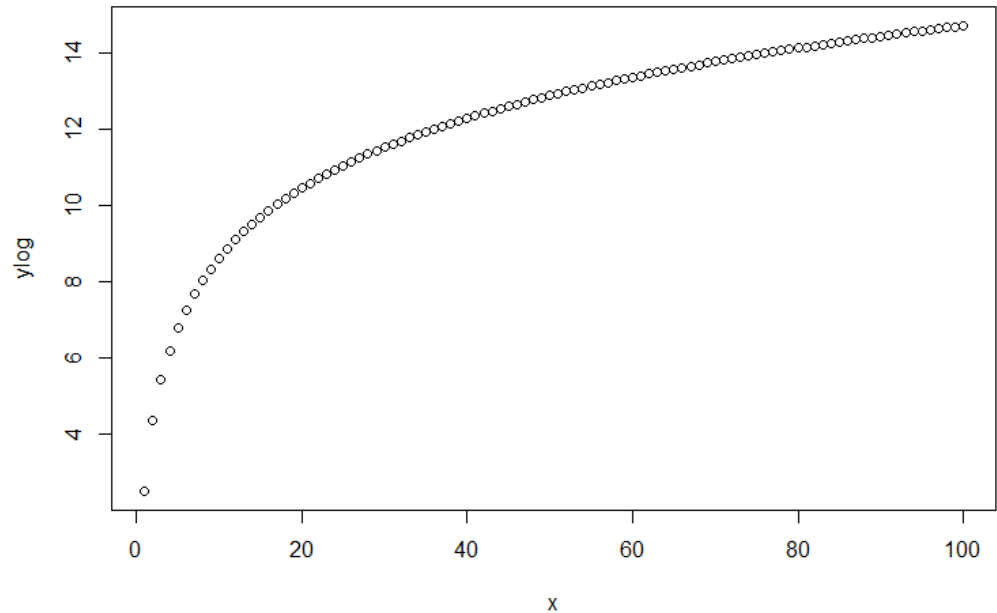
```
ylin <- 2.5 + 2.65*x
```

```
plot(x,ylin)
```



```
ylog <- 2.5 + 2.65*log(x)
```

```
plot(x,ylog)
```



Parametric response functions

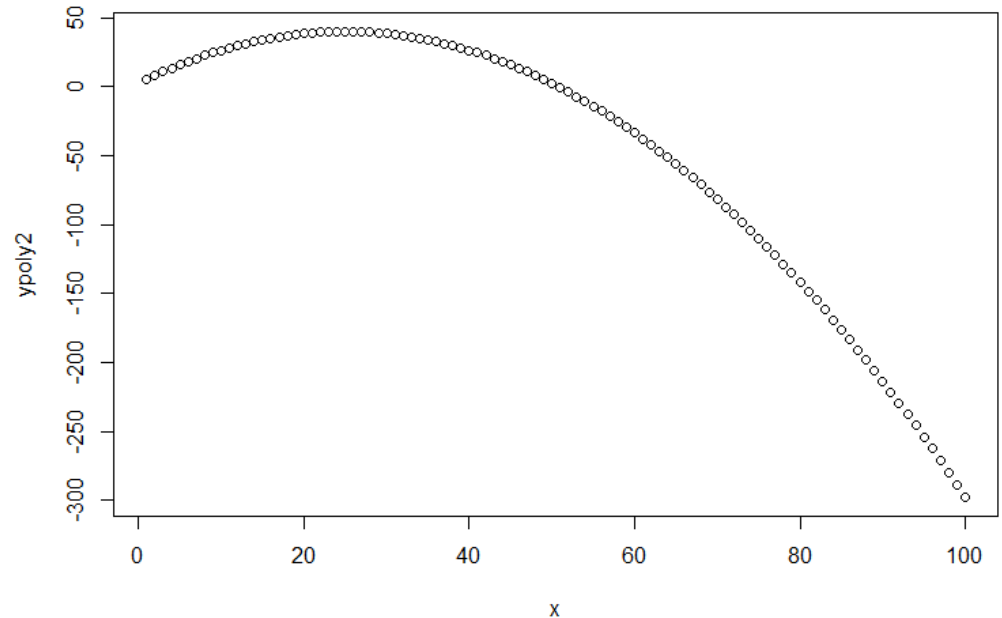
```
# linear parametric functions
```

```
x <- seq(1,100,1)
```

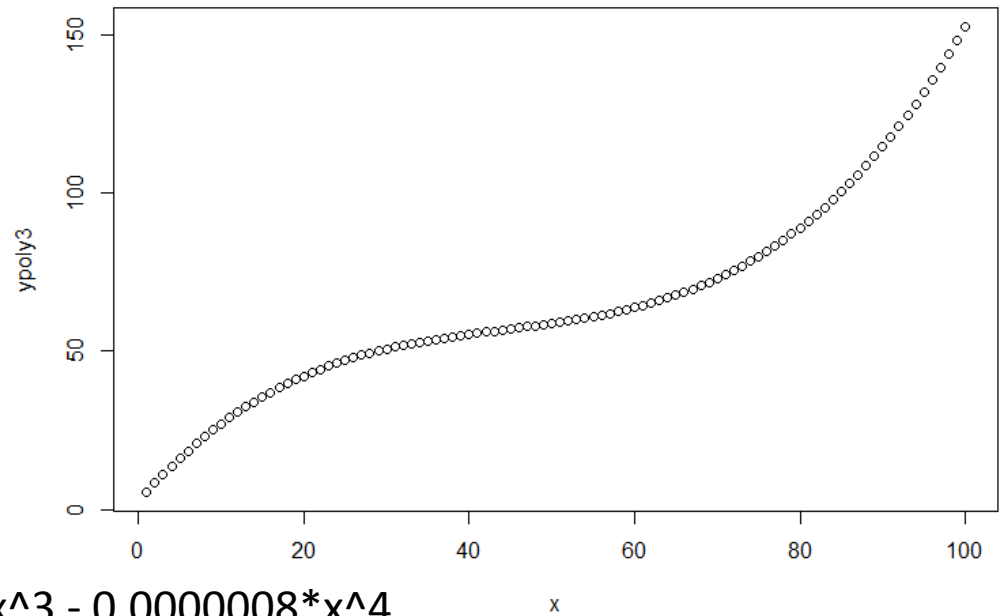
```
# Quadratic, second order polynomial
```

```
ypoly2 <- 2.5 + 3*x - 0.06*x^2
```

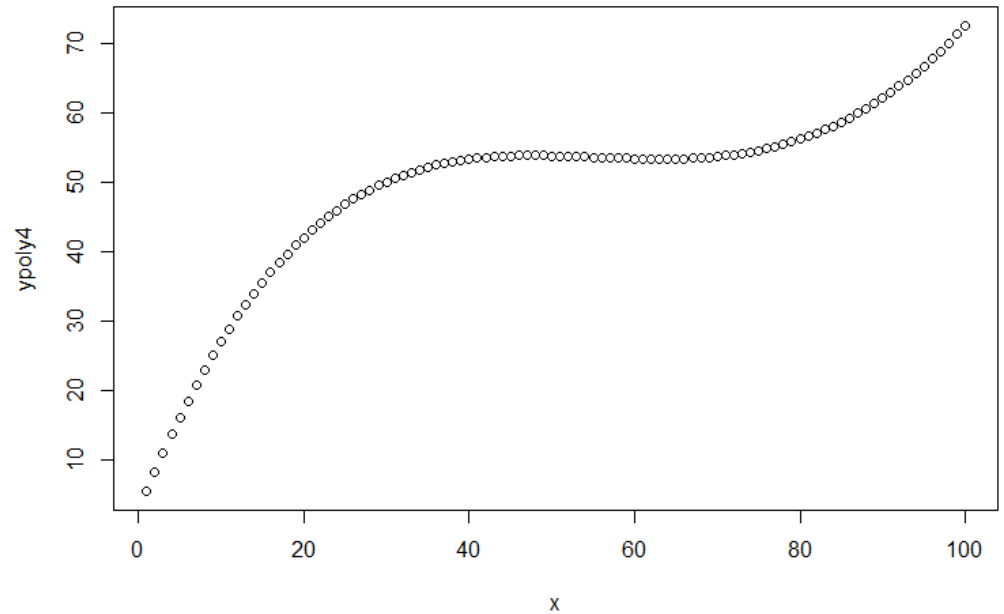
```
plot(x,ypoly2)
```



```
ypoly3 <- 2.5 + 3*x - 0.06*x^2 + 0.00045*x^3  
plot(x,ypoly3) # cubic, third order polynomial
```



```
ypoly4 <- 2.5 + 3*x - 0.06*x^2 + 0.00045*x^3 - 0.0000008*x^4  
plot(x,ypoly4)
```

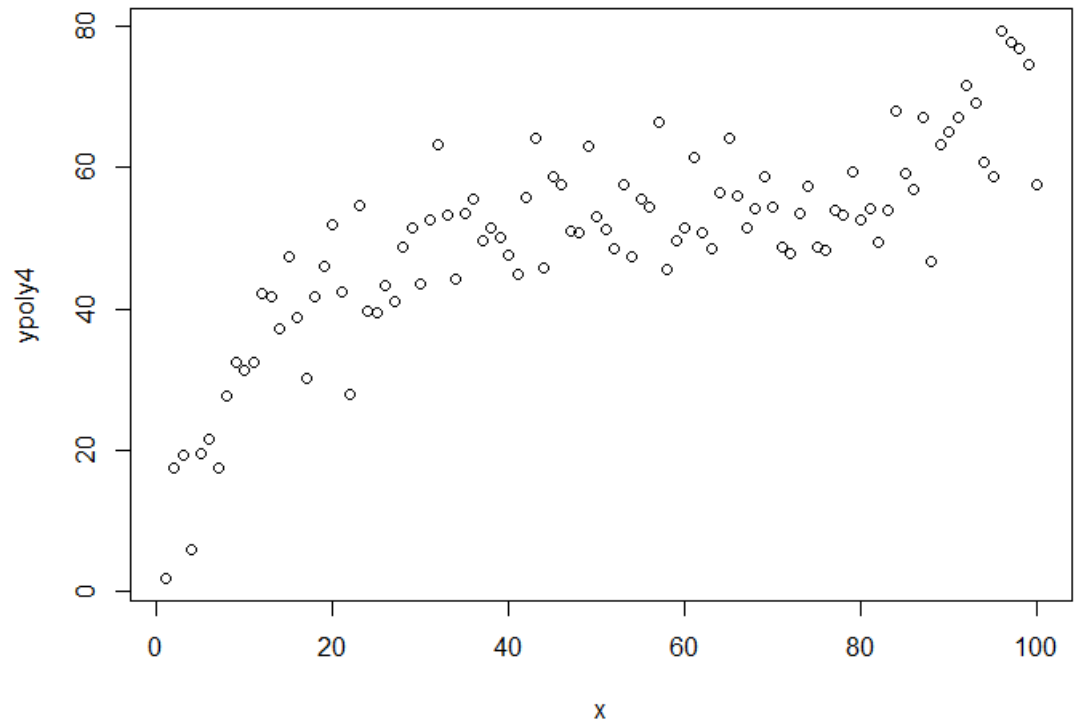


No a-priori relationship between x and y

Non-parametric smoothed functions

```
ypoly4 <- 2.5 + 3*x - 0.06*x^2 + 0.00045*x^3 - 0.0000008*x^4 + rnorm(length(x),0,6)  
plot(x,ypoly4)
```

Let the data
determine the
shape of the
response curve



Non-parametric smoothed functions

```
ypoly4 <- 2.5 + 3*x - 0.06*x^2 + 0.00045*x^3 - 0.0000008*x^4 + rnorm(length(x),0,6)
```

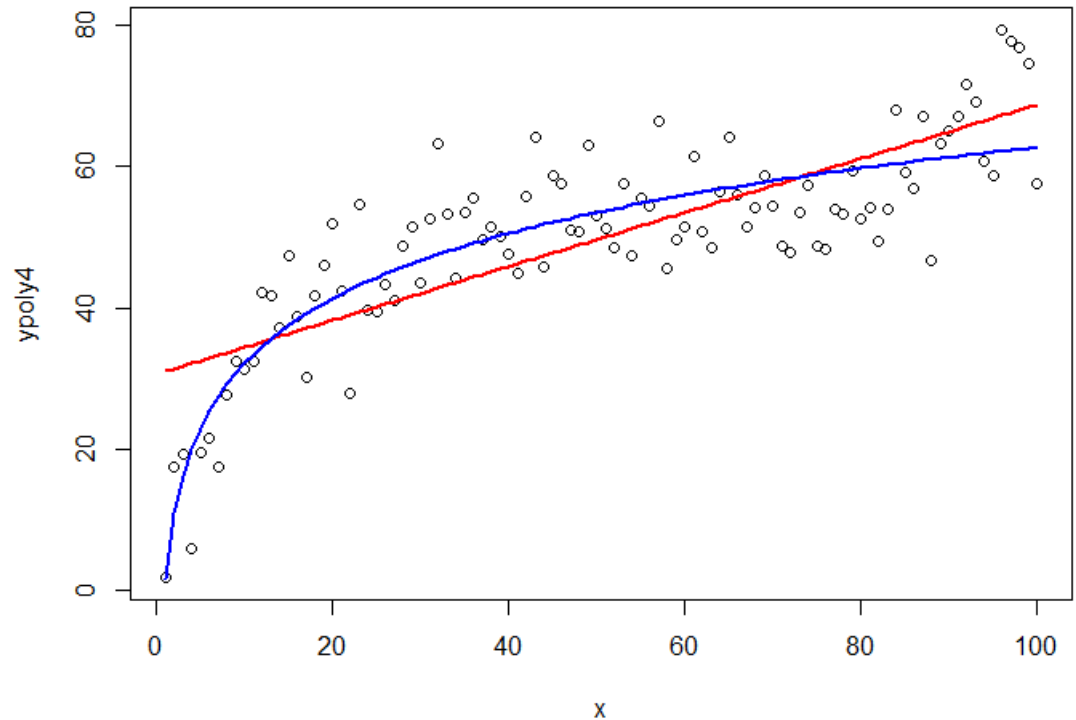
```
ylm1 <- lm(ypoly4 ~ x)
```

```
ylm2 <- lm(ypoly4 ~ log(x))
```

```
plot(x,ypoly4)
```

```
lines(x,ylm1$fitted.values,col="red",lwd=2)
```

```
lines(x,ylm2$fitted.values,col="blue",lwd=2)
```



Non-parametric smoothed functions

```
ypoly4 <- 2.5 + 3*x - 0.06*x^2 + 0.00045*x^3 - 0.0000008*x^4 + rnorm(length(x),0,6)
```

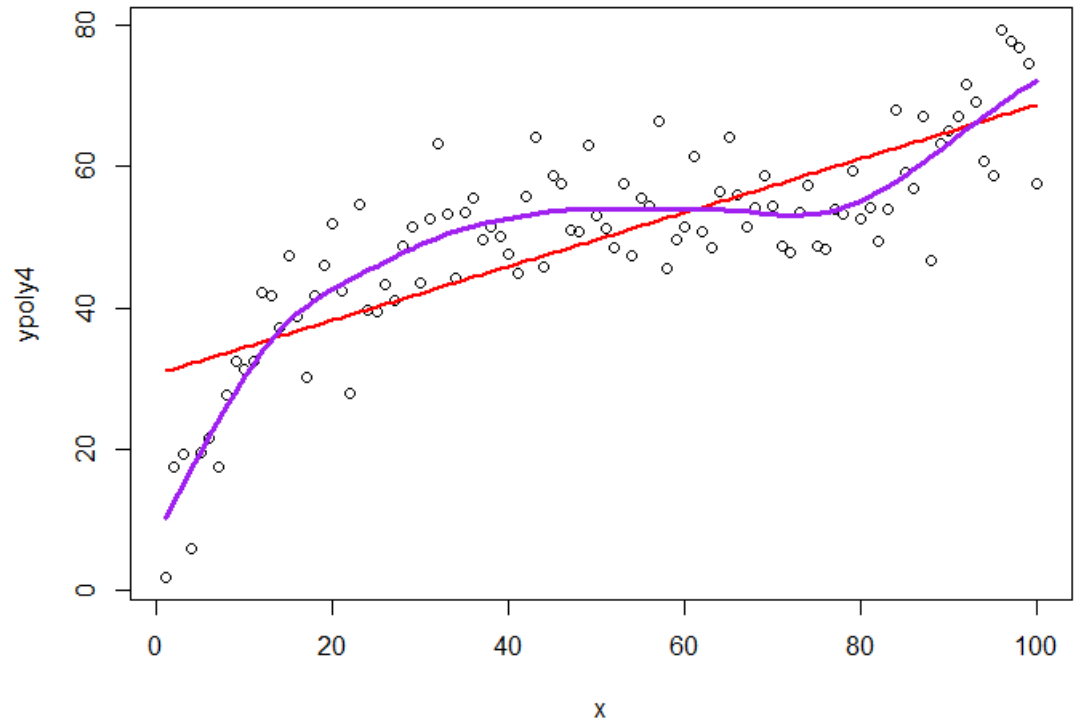
```
ylm1 <- lm(ypoly4 ~ x)
```

```
spline1 <- smooth.spline(x,ypoly4)
```

```
plot(x,ypoly4)
```

```
lines(x,ylm1$fitted.values,col="red",lwd=2)
```

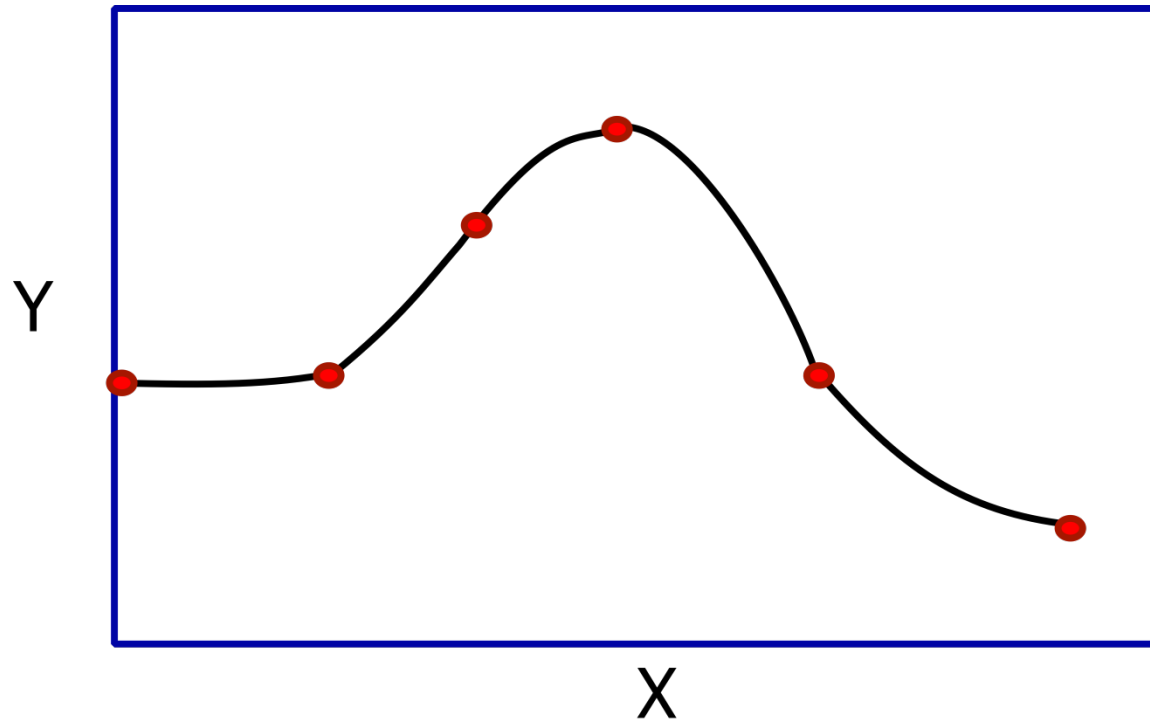
```
lines(predict(spline1,x),col="purple",lwd=3)
```



Non-parametric smoothed functions

- Spline
 - A numeric function that is **piecewise-defined** by **polynomial functions**, and which possesses a sufficiently high degree of **smoothness** at the places where the polynomial pieces connect (knots).

Spline

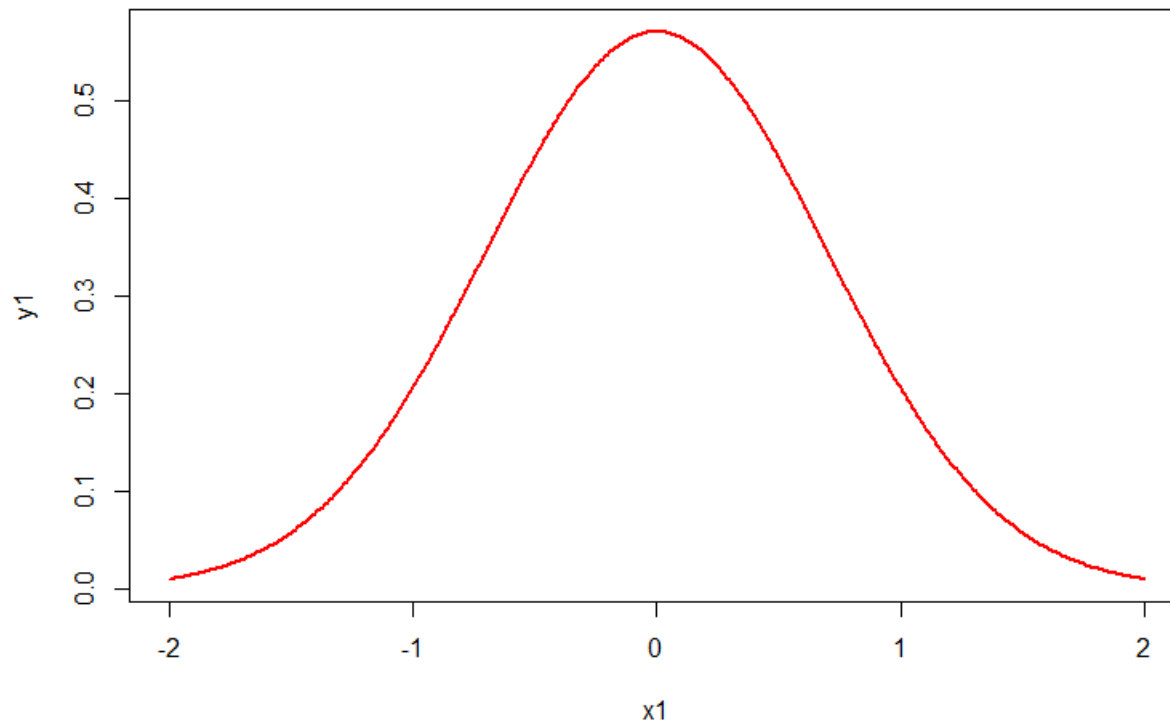


• Knots
polynomial function

cubic splines: third order polynomials

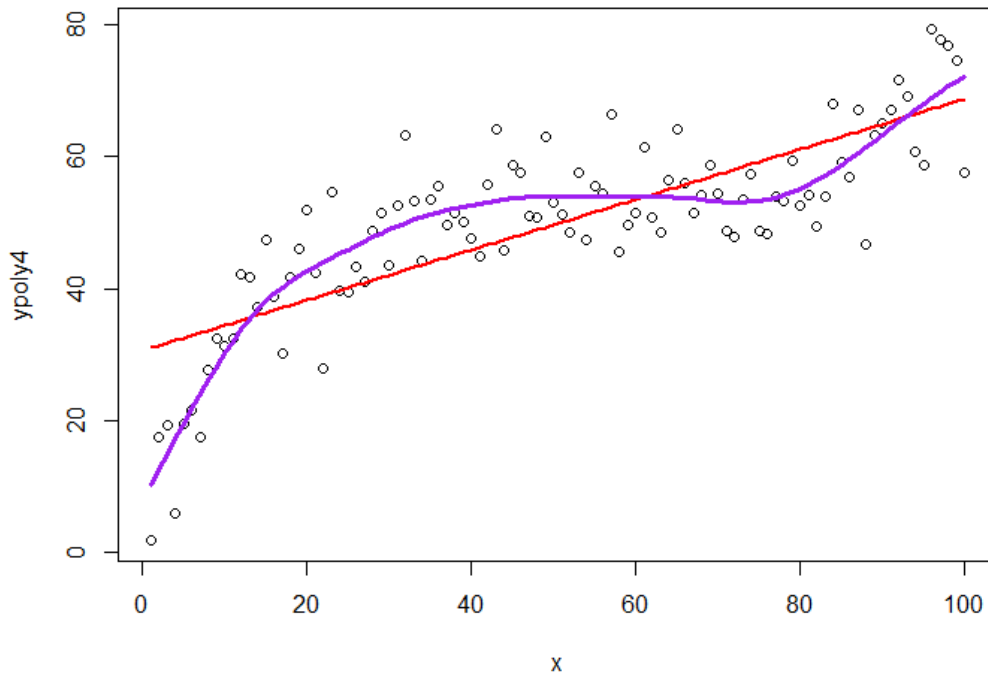
Splines

$$f_X(x) = \begin{cases} \frac{1}{4}(x+2)^3 & -2 \leq x \leq -1 \\ \frac{1}{4}(3|x|^3 - 6x^2 + 4) & -1 \leq x \leq 1 \\ \frac{1}{4}(2-x)^3 & 1 \leq x \leq 2 \end{cases}$$



Overfitting (noise vs. process)?

- Penalizing badness of fit?
- Penalizing wiggleness?



Generalized additive models

- Linear model:
 - Linear combination of parametric terms predict response variable
- Generalized linear model:
 - Parametric predictor terms related to a response variable through a linker function
- Generalized additive model:
 - A GAM is a generalized linear model in which the linear predictor is given by a user specified sum of smooth functions of the covariates plus a conventional parametric component of the linear predictor

lm, glm, gam

$$y_i = B_0 + B_1x_i + \cdots + B_pX_p + \varepsilon_i$$

$$g(y_i) = n_i = B_0 + B_1x_i + \cdots + B_pX_p + \varepsilon_i$$

$$g(y_i) = n_i = B_0 + f_1(x_{1i}) + \cdots + f_p(x_{1p}) + \varepsilon_i$$

GAM is a GLM where the linear predictor depends on smooth functions of covariates

GAM specification

```
install.packages("mgcv")  
library(mgcv)
```

$$y \sim s(x)$$

Single non-parametrically
smoothed function

$$y \sim s(x) + s(w) + s(z)$$

Multiple smoothed functions

$$y \sim s + s(w)$$

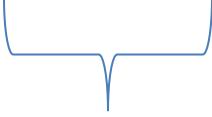
Mixture of parametric parameters
and smoothed function

$$y \sim s(x) + s(z) + s(x, z)$$

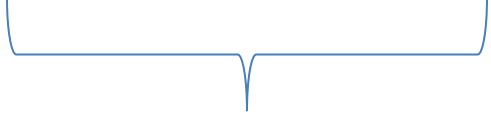
Inclusion of nested smoothed terms
(two-dimensional)

GAM specification

```
gm <- gam(growth ~ s(temp) + s(precip), family=gaussian())
```



Non-parametric
smoothed function



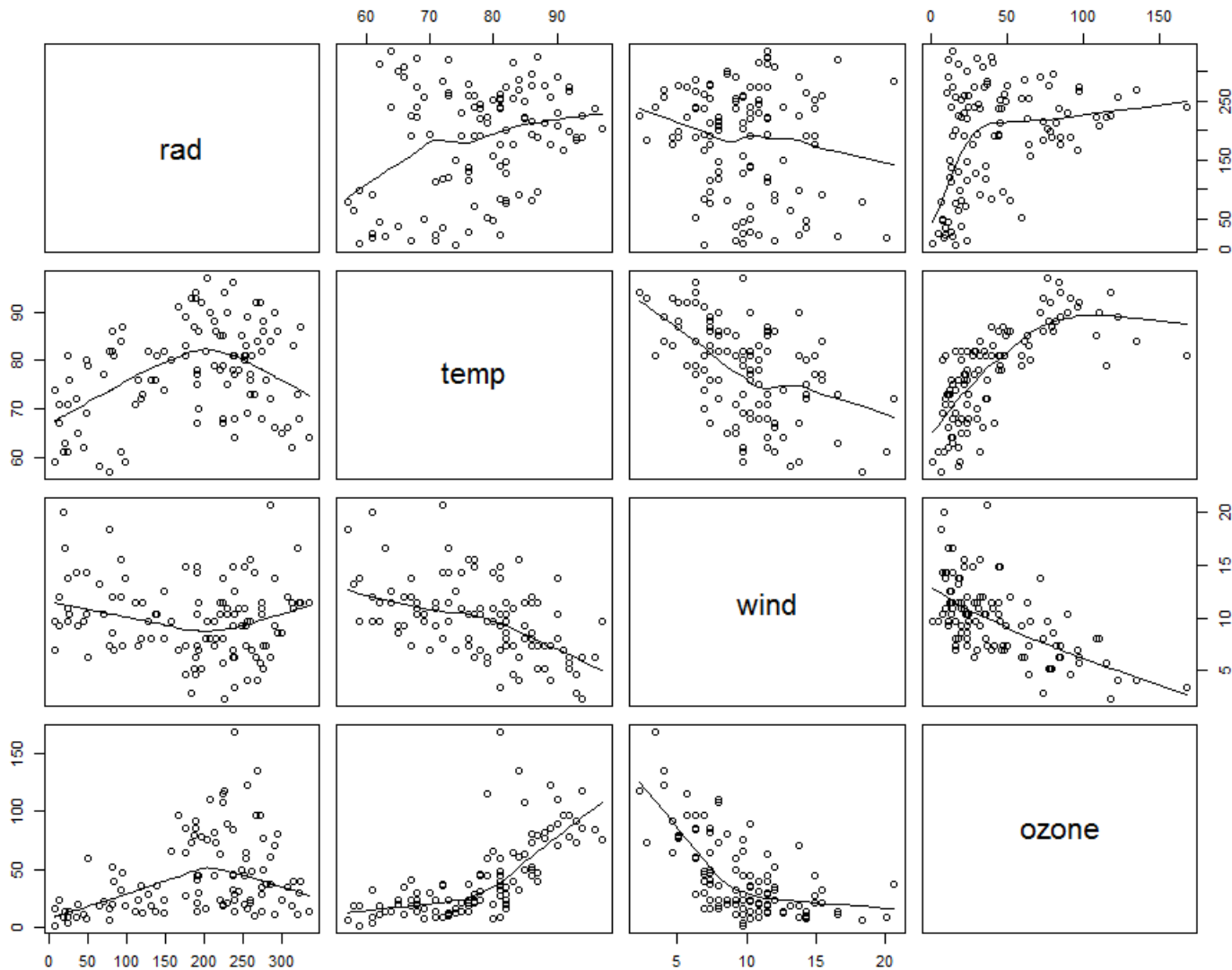
Linker function

GAM example 1

- Local ozone concentration

rad	temp	wind	ozone
Min. : 7.0	Min. :57.00	Min. : 2.300	Min. : 1.0
1st Qu.:113.5	1st Qu.:71.00	1st Qu.: 7.400	1st Qu.: 18.0
Median :207.0	Median :79.00	Median : 9.700	Median : 31.0
Mean :184.8	Mean :77.79	Mean : 9.939	Mean : 42.1
3rd Qu.:255.5	3rd Qu.:84.50	3rd Qu.:11.500	3rd Qu.: 62.0
Max. :334.0	Max. :97.00	Max. :20.700	Max. :168.0

Local ozone concentration



Local ozone concentration

```
model <- gam(ozone ~ s(rad) + s(temp) + s(wind))
summary(model)
```

Family: gaussian

Link function: identity

Formula:

ozone ~ s(rad) + s(temp) + s(wind)

Parametric coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	42.10	1.66	25.36	<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(rad)	2.763	3.451	3.823	0.00931 **
s(temp)	3.841	4.762	11.767	7.18e-09 ***
s(wind)	2.918	3.666	13.770	1.46e-08 ***

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

R-sq.(adj) = 0.724 Deviance explained = 74.8%

GCV = 338 Scale est. = 305.96 n = 111

Local ozone concentration

```
plot(model)
```

Temp

Min. :57.00

1st Qu.:71.00

Median :79.00

Mean :77.79

3rd Qu.:84.50

Max. :97.00

wind

Min. : 2.300

1st Qu.: 7.400

Median : 9.700

Mean : 9.939

3rd Qu.:11.500

Max. :20.700

ozone

Min. : 1.0

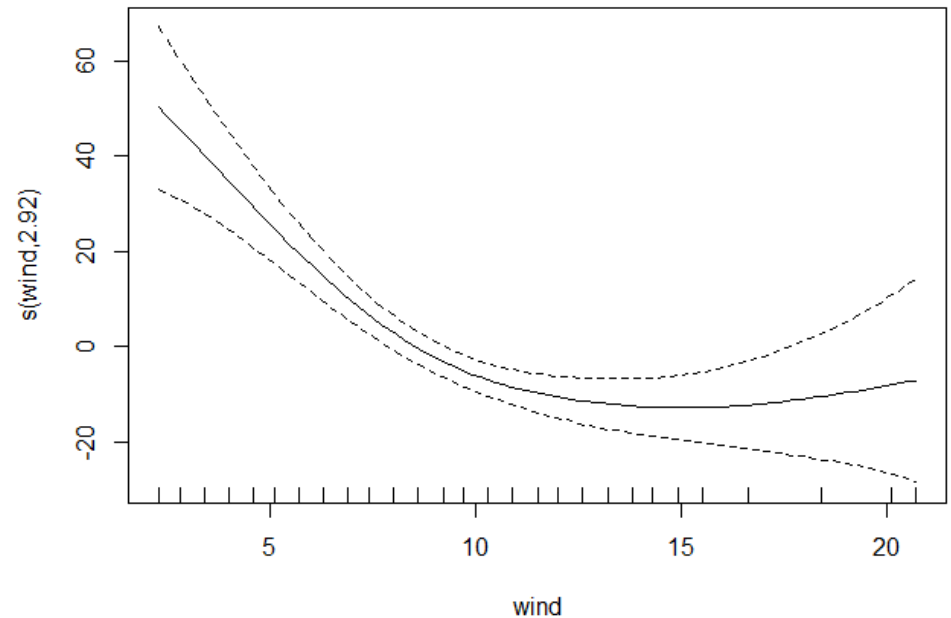
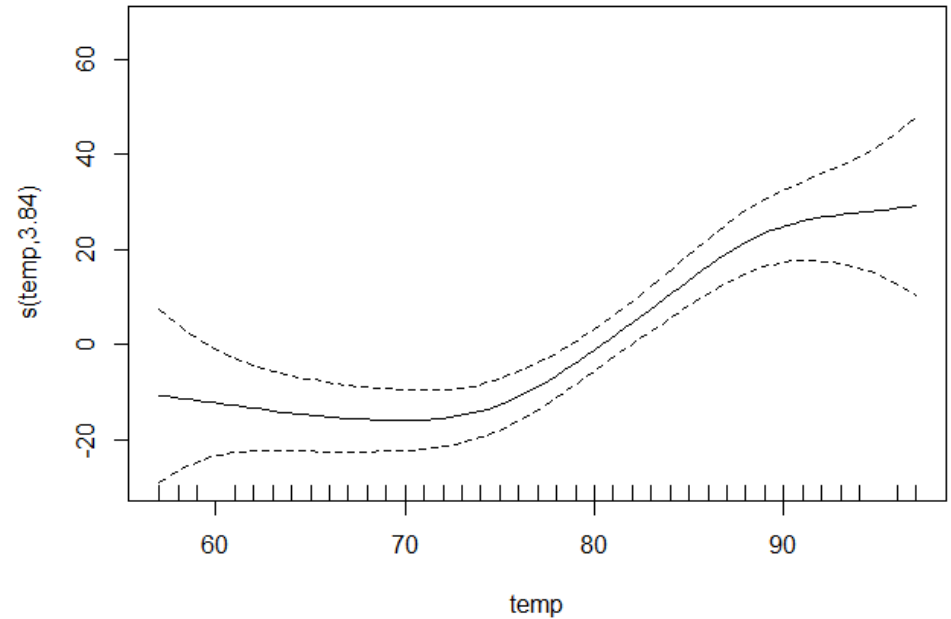
1st Qu.: 18.0

Median : 31.0

Mean : 42.1

3rd Qu.: 62.0

Max. :168.0



GAM example 2 (binary data)



Presence
Absence

incidence
Min. :0.00
1st Qu.:0.00
Median :1.00
Mean :0.58
3rd Qu.:1.00
Max. :1.00

area
Min. :0.153
1st Qu.:2.248
Median :4.170
Mean :4.319
3rd Qu.:6.431
Max. :9.269

isolation
Min. :2.023
1st Qu.:4.823
Median :5.801
Mean :5.856
3rd Qu.:7.191
Max. :9.577



Species occurrence

```
model <- gam(incidence ~ s(area) + s(isolation), family = binomial)
summary(model)
```

Family: binomial

Link function: logit

Formula:

incidence ~ s(area) + s(isolation)

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	1.6371	0.9898	1.654	0.0981 .

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

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value
s(area)	2.429	3.066	3.455	0.33606
s(isolation)	1.000	1.000	7.480	0.00624 **

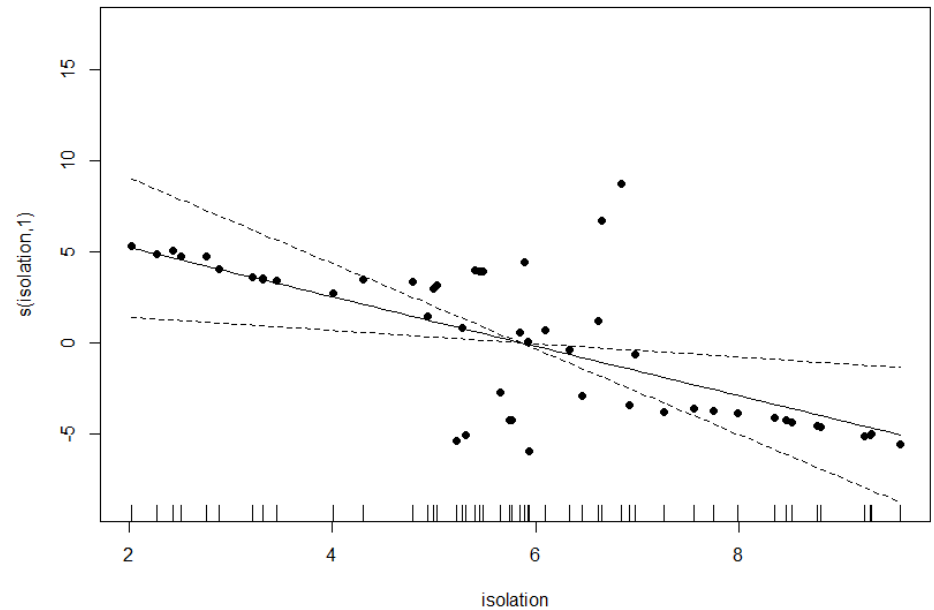
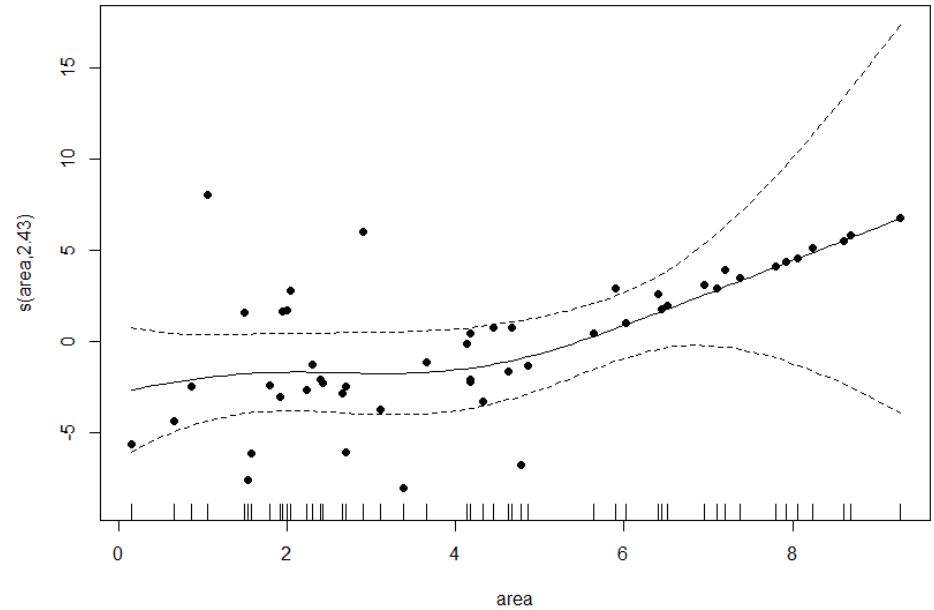
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) = 0.63 Deviance explained = 63.1%

UBRE = -0.32096 Scale est. = 1 n = 50

Species occurrence

```
plot.gam(model,residuals=T,pch=16)
```



Species occurrence

```
model1 <- gam(incidence~s(area) + isolation,family=binomial)
summary(model1)
```

Family: binomial
Link function: logit

Formula:
incidence ~ s(area) + isolation

Parametric coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	9.5755	3.1859	3.006	0.00265 **
isolation	-1.3555	0.4956	-2.735	0.00624 **

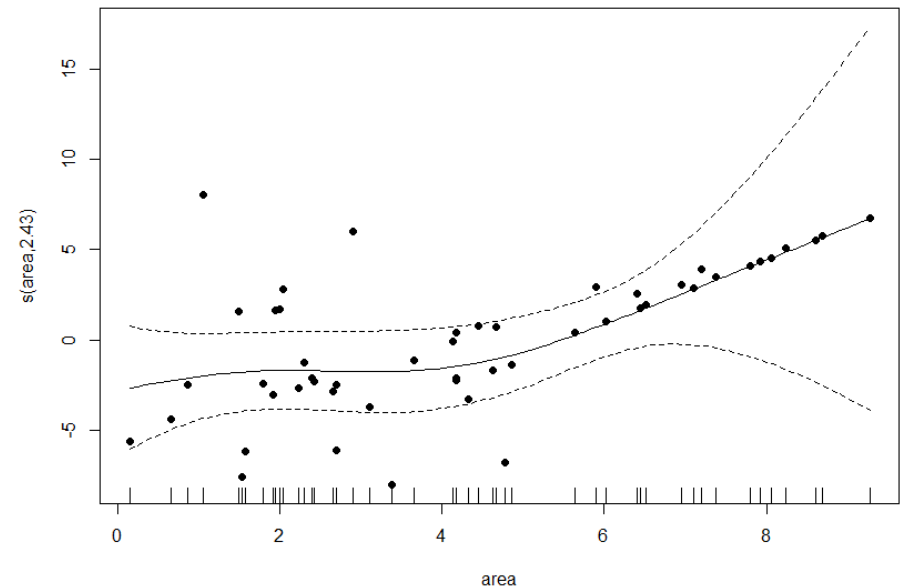
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Approximate significance of smooth terms:

	edf	Ref.df	Chi.sq	p-value
s(area)	2.429	3.066	3.555	0.323

R-sq.(adj) = 0.63 Deviance explained = 63.1%

UBRE = -0.32096 Scale est. = 1 n = 50



GAM vs. GLM

- GAM
 - Good
 - Extremely flexible models for fitting smooth curves
 - No a-priori assumptions of response curve shape
 - With ecological data the fit of GAM is often better than GLM or LM
 - Bad
 - Hard to evaluate (except graphically)
 - Can't provide an equation for a paper
 - Hard (impossible) to interpret and understand
 - Biological and physical processes underlying response curve
 - Computationally intensive (not an issue normally)
 - Statistical power vs. ecological interpretability

GAM details

Wood S.N. (2006)

Generalized Additive Models: An Introduction with R.
Chapman and Hall/CRC Press