

# Moving Beyond Linearity

by

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## Polynomial Regression:

The nonlinear relationship between the predictors and the response is defined by a polynomial function

$$y_i = \beta_0 + \beta_1 x_i + \beta_2 x_i^2 + \beta_3 x_i^3 + \cdots + \beta_d x_i^d + \epsilon_i$$

where  $\epsilon_i$  is the error term.

- Polynomial regression extends the linear model by adding extra predictors, obtained by raising each of the original predictors to a power.
- Least squares returns variance estimates for each of the fitted coefficients  $\beta_j^2$  as well as the covariances between pairs of coefficient estimates.

```
library (ISLR2)
attach (Wage)
fit <- lm(wage ~ poly (age , 4), data = Wage)
coef ( summary (fit))

##              Estimate Std. Error    t value    Pr(>|t|)
## (Intercept)   111.70361   0.7287409  153.283015 0.000000e+00
## poly(age, 4)1  447.06785  39.9147851   11.200558 1.484604e-28
## poly(age, 4)2 -478.31581  39.9147851  -11.983424 2.355831e-32
## poly(age, 4)3  125.52169  39.9147851    3.144742 1.678622e-03
## poly(age, 4)4  -77.91118  39.9147851   -1.951938 5.103865e-02

fit2 <- lm(wage ~ poly (age , 4, raw = T), data = Wage)
coef ( summary (fit2))

##              Estimate    Std. Error    t value    Pr(>|t|)
## (Intercept)   -1.841542e+02  6.004038e+01  -3.067172 0.0021802539
## poly(age, 4, raw = T)1  2.124552e+01  5.886748e+00   3.609042 0.0003123618
## poly(age, 4, raw = T)2 -5.638593e-01  2.061083e-01  -2.735743 0.0062606446
## poly(age, 4, raw = T)3  6.810688e-03  3.065931e-03   2.221409 0.0263977518
## poly(age, 4, raw = T)4 -3.203830e-05  1.641359e-05  -1.951938 0.0510386498

fit2a <- lm(wage ~ age + I(age^2) + I(age^3) + I(age^4), data = Wage)
coef (fit2a)
```

```
##      (Intercept)          age      I(age^2)      I(age^3)      I(age^4)
## -1.841542e+02  2.124552e+01 -5.638593e-01  6.810688e-03 -3.203830e-05

fit2b <- lm(wage ~ cbind (age , age^2, age^3, age^4), data = Wage)
agelims <- range (age)
age.grid <- seq (from = agelims[1], to = agelims [2])
preds <- predict (fit , newdata = list (age = age.grid), se = TRUE)
se.bands <- cbind (preds$fit + 2 * preds$se.fit , preds$fit - 2 * preds$se.fi
t)
library(ggplot2)
ggplot()+
  geom_point(aes(Wage$age , Wage$wage),col="pink")+
  geom_smooth(aes(Wage$age , Wage$wage),col="red")

## `geom_smooth()` using method = 'gam' and formula = 'y ~ s(x, bs = "cs")'
```



```
labs(title = "Degree -4 Polynomial" , x= "age" , y="Wage")

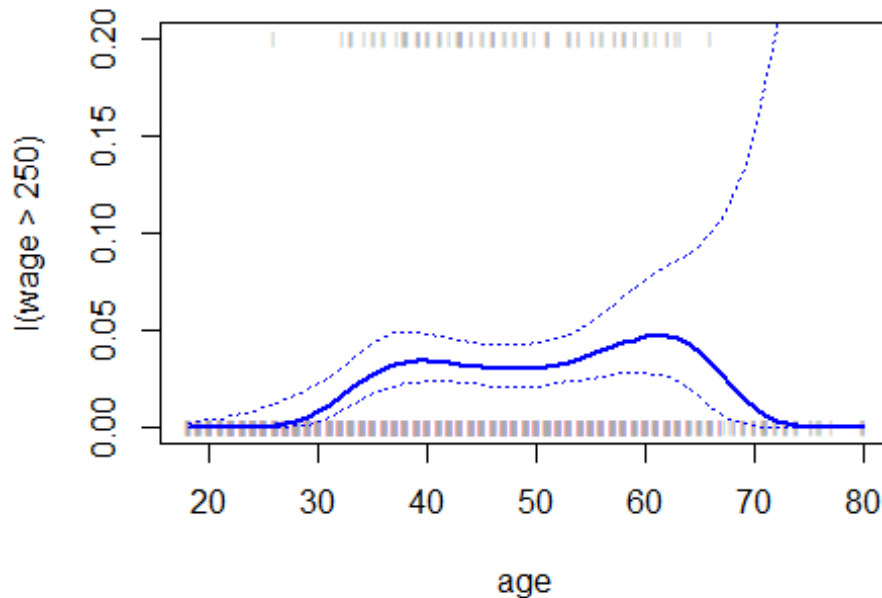
## $x
## [1] "age"
##
## $y
## [1] "Wage"
##
## $title
## [1] "Degree -4 Polynomial"
##
```

```
## attr(,"class")
## [1] "labels"

preds2 <- predict (fit2 , newdata = list (age = age.grid), se = TRUE)
max (abs (preds$fit - preds2$fit))

## [1] 7.81597e-11

fit <- glm (I(wage > 250) ~ poly (age , 4), data = Wage , family = binomial)
preds <- predict (fit , newdata = list (age = age.grid), se = T)
pfit <- exp (preds$fit) / (1 + exp (preds$fit))
se.bands.logit <- cbind (preds$fit + 2 * preds$se.fit ,
preds$fit - 2 * preds$se.fit)
se.bands <- exp (se.bands.logit) / (1 + exp (se.bands.logit))
preds <- predict (fit , newdata = list (age = age.grid), type = "response", se = T)
plot (age , I(wage > 250), xlim = agelims , type = "n", ylim = c(0, .2))
points ( jitter (age), I((wage > 250) / 5), cex = .5, pch = "|", col = "darkgrey")
lines (age.grid, pfit , lwd = 2, col = "blue")
matlines (age.grid , se.bands , lwd = 1, col = "blue", lty = 3)
```



```
table ( cut (age , 4))

##
## (17.9,33.5] (33.5,49] (49,64.5] (64.5,80.1]
##          750      1399      779      72
```

```
fit <- lm(wage ~ cut (age , 4), data = Wage)
coef(summary (fit))
```

##	Estimate	Std. Error	t value	Pr(> t )
## (Intercept)	94.158392	1.476069	63.789970	0.000000e+00
## cut(age, 4)(33.5,49]	24.053491	1.829431	13.148074	1.982315e-38
## cut(age, 4)(49,64.5]	23.664559	2.067958	11.443444	1.040750e-29
## cut(age, 4)(64.5,80.1]	7.640592	4.987424	1.531972	1.256350e-01

### Step Functions:

Step functions cut the range of a variable into  $K$  distinct regions in order to produce a qualitative variable. We create cutpoints  $c_1, c_2, \dots, c_K$  in the range of  $X$ , and then construct  $K + 1$  new variables.

$$\begin{aligned}
 C_0(X) &= I(X < c_1) \\
 C_1(X) &= I(c_1 \leq X < c_2) \\
 C_2(X) &= I(c_2 \leq X < c_3) \\
 &\vdots \\
 C_{K-1}(X) &= I(c_{K-1} \leq X < c_K) \\
 C_K(X) &= I(c_K \leq X)
 \end{aligned}$$

We then use least squares to fit the linear model

$$y_i = \beta_0 + \beta_1 C_1(x_i) + \beta_2 C_2(x_i) + \beta_3 C_3(x_i) + \dots + \beta_K C_K(x_i) + \epsilon_i$$

### Regression Splines:

- Regression splines are more flexible than polynomials and step functions, and in fact are an extension of the two. They involve dividing the range of  $X$  into  $K$  distinct regions. Within each region, a polynomial function is fit to the data. However, these polynomials are constrained so that they join smoothly at the region boundaries, or knots.

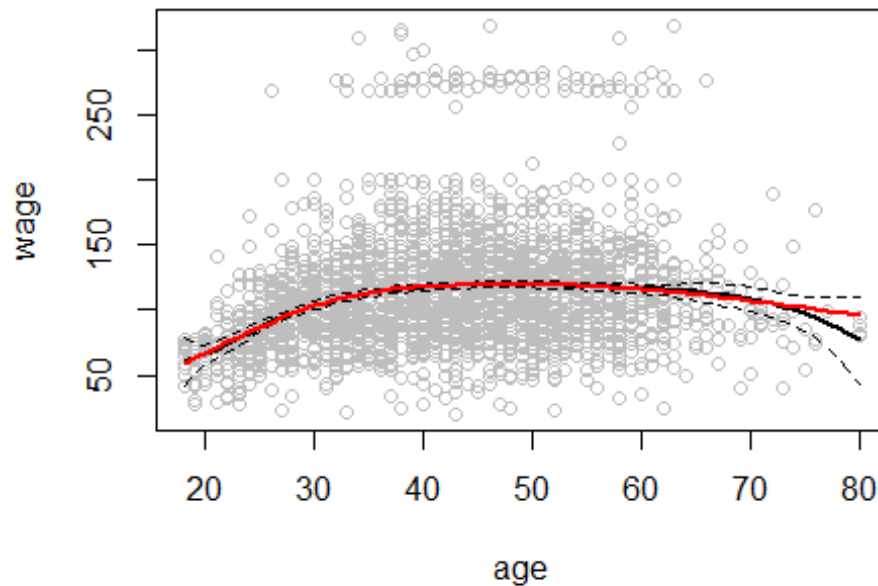
A piecewise cubic polynomial with a single knot at a point  $c$  takes the form

$$y_i = \begin{cases} \beta_{01} + \beta_{11}x_i + \beta_{21}x_i^2 + \beta_{31}x_i^3 + \epsilon_i & \text{if } x_i < c \\ \beta_{02} + \beta_{12}x_i + \beta_{22}x_i^2 + \beta_{32}x_i^3 + \epsilon_i & \text{if } x_i \geq c. \end{cases}$$

- To choose number of knots an approach is to use cross-validation. With this method, we remove a portion of the data (say 10 %), fit a spline with a certain number of knots to the remaining data and then use the spline to make predictions for the held-out portion.

```
library (splines)
fit <- lm(wage ~ bs(age , knots = c(25, 40, 60)), data = Wage)
pred <- predict (fit , newdata = list (age = age.grid), se = T)
plot (age , wage , col = "gray")
lines (age.grid, pred$fit , lwd = 2)
lines (age.grid , pred$fit + 2 * pred$se, lty = "dashed")
lines (age.grid , pred$fit - 2 * pred$se, lty = "dashed")
fit2 <- lm(wage ~ ns(age , df = 4), data = Wage)
```

```
pred2 <- predict (fit2 , newdata = list (age = age.grid), se = T)
lines (age.grid , pred2$fit , col = "red", lwd = 2)
```



```
plot (age , wage , xlim = agelims , cex = .5, col = "darkgrey")
title ("Smoothing Spline")
fit <- smooth.spline (age , wage , df = 16)
fit2 <- smooth.spline (age , wage , cv = TRUE)

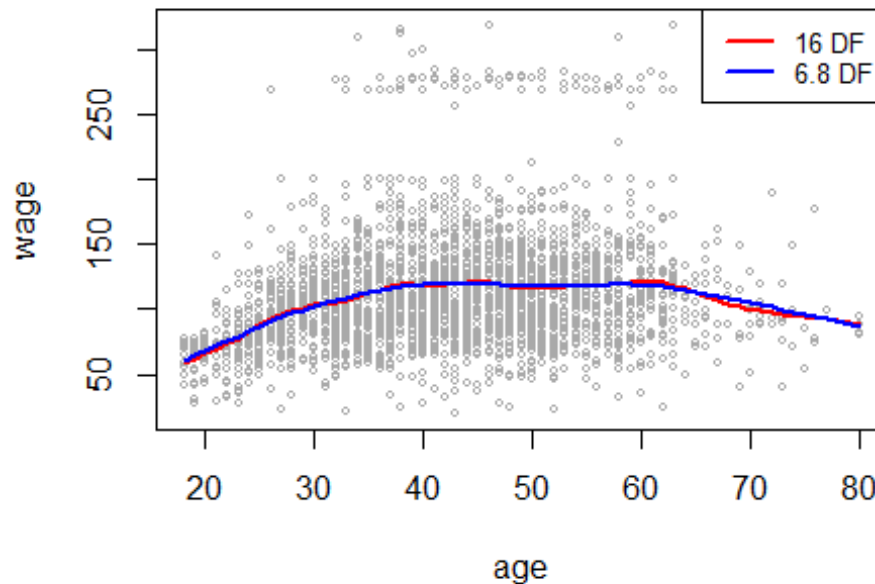
## Warning in smooth.spline(age, wage, cv = TRUE): cross-validation with non-
## unique
## 'x' values seems doubtful

fit2$df

## [1] 6.794596

lines (fit , col = "red", lwd = 2)
lines (fit2 , col = "blue", lwd = 2)
legend ("topright", legend = c("16 DF", "6.8 DF") , col = c("red", "blue"), l
ty = 1, lwd = 2, cex = .8)
```

## Smoothing Spline



### Smoothing Splines:

Smoothing splines result from minimizing a residual sum of squares criterion subject to a smoothness penalty. We really want is a function  $g$  that makes RSS small but that is also smooth. A natural approach is to find the function  $g$  that minimizes

$$\sum_{i=1}^n (y_i - g(x_i))^2 + \lambda \int g''(t)^2 dt$$

where  $\lambda$  is a nonnegative tuning parameter and chosen using LOOCV. The equation takes the “Loss+Penalty” formulation.

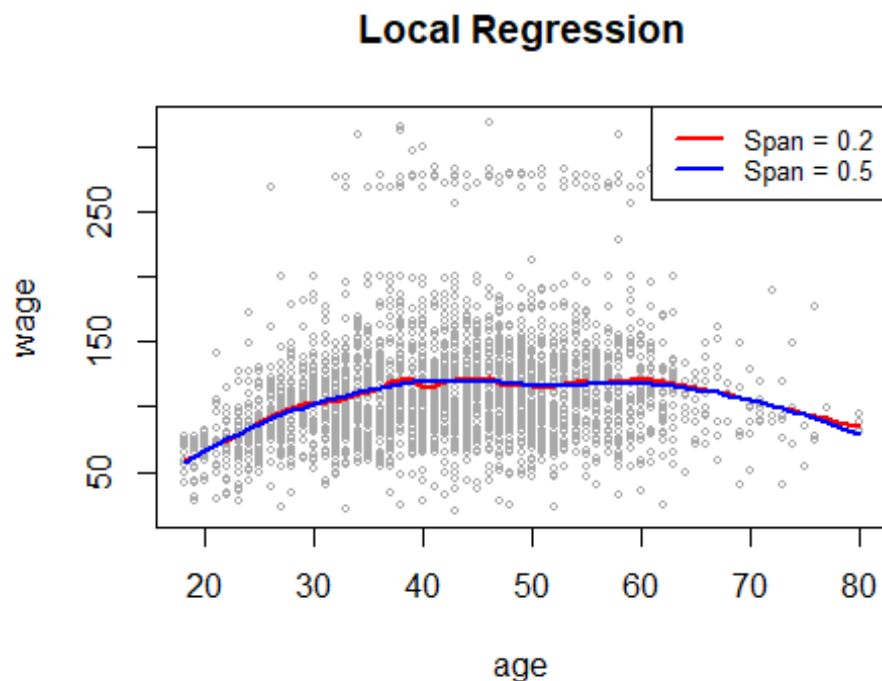
### Local Regression:

- In Local regression the regions are allowed to overlap, and indeed they do so in a very smooth way.
1. Gather the fraction  $s = k/n$  of training points whose  $x_i$  are closest to  $x_0$ .
  2. Assign a weight  $K_{i0} = K(x_i, x_0)$  to each point in this neighborhood, so that the point furthest from  $x_0$  has weight zero, and the closest has the highest weight. All but these  $k$  nearest neighbors get weight zero.
  3. Fit a weighted least squares regression of the  $y_i$  on the  $x_i$  using the aforementioned weights, by finding  $\hat{\beta}_0$  and  $\hat{\beta}_1$  that minimize  $\sum_{i=1}^n K_{i0} (y_i - \beta_0 - \beta_1 x_i)^2$ .
  4. The fitted value at  $x_0$  is given by  $\hat{f}(x_0) = \hat{\beta}_0 + \hat{\beta}_1 x_0$

```

plot (age , wage , xlim = agelims , cex = .5, col = "darkgrey")
title ("Local Regression")
fit <- loess (wage ~ age , span = .2, data = Wage)
fit2 <- loess (wage ~ age , span = .5, data = Wage)
lines (age.grid, predict (fit , data.frame (age = age.grid)),
col = "red", lwd = 2)
lines (age.grid, predict (fit2 , data.frame (age = age.grid)),
col = "blue", lwd = 2)
legend ("topright", legend = c("Span = 0.2", "Span = 0.5") ,
col = c("red", "blue"), lty = 1, lwd = 2, cex = .8)

```



### Generalized Additive Models:

Generalized additive models (GAMs) provide a general framework for extending a standard linear model by allowing non-linear functions of each of the variables, while maintaining additivity.

```

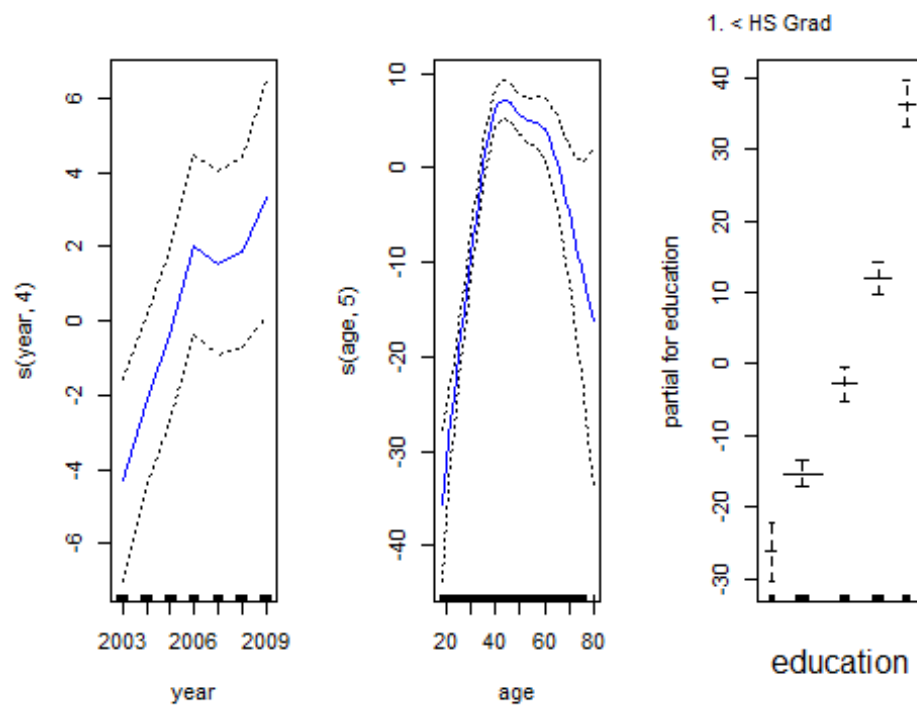
gam1 <- lm(wage ~ ns(year , 4) + ns(age , 5) + education , data = Wage)
library (gam)

## Loading required package: foreach

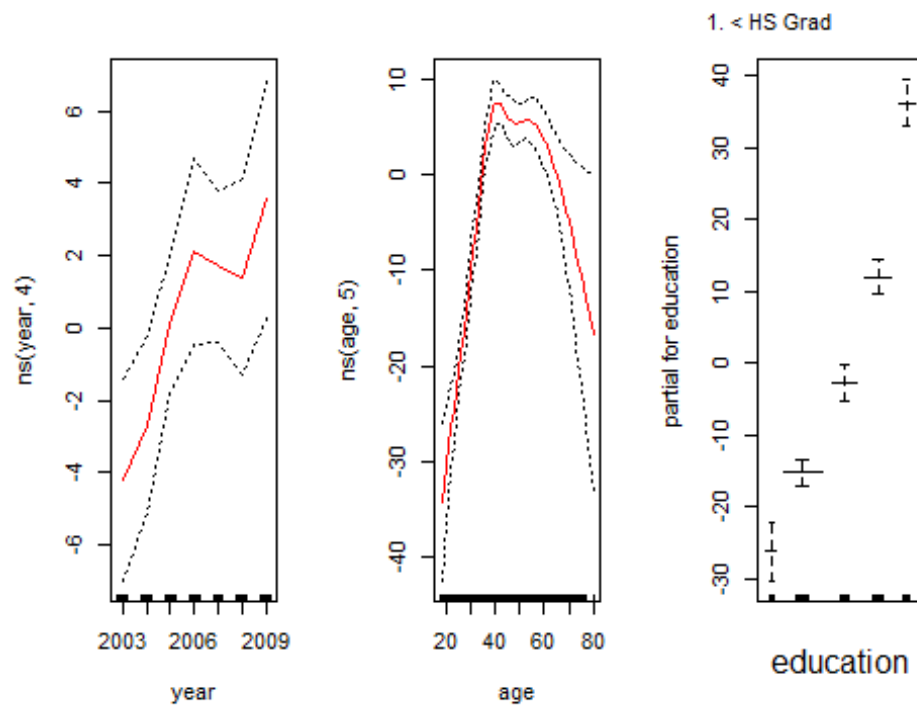
## Loaded gam 1.22

gam.m3 <- gam (wage ~ s(year , 4) + s(age , 5) + education , data = Wage)
par (mfrow = c(1, 3))
plot (gam.m3, se = TRUE , col = "blue")

```



```
plot.Gam (gam1 , se = TRUE , col = "red")
```





```

gam.m1 <- gam (wage ~ s(age , 5) + education , data = Wage)
gam.m2 <- gam (wage ~ year + s(age , 5) + education , data = Wage)
anova (gam.m1 , gam.m2 , gam.m3 , test = "F")

## Analysis of Deviance Table
##
## Model 1: wage ~ s(age, 5) + education
## Model 2: wage ~ year + s(age, 5) + education
## Model 3: wage ~ s(year, 4) + s(age, 5) + education
##   Resid. Df Resid. Dev Df Deviance      F    Pr(>F)
## 1      2990      3711731
## 2      2989      3693842  1  17889.2 14.4771 0.0001447 ***
## 3      2986      3689770  3   4071.1  1.0982 0.3485661
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```