# Untitled

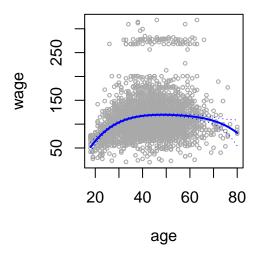
```
library(ISLR2)
attach(Wage)
library(gam)
Loading required package: splines
Loading required package: foreach
Loaded gam 1.22-5
library(glmnet)
Loading required package: Matrix
Loaded glmnet 4.1-8
library(boot)
library(ggplot2)
library(leaps)
#labs #7.8.1 Polynomial Regression and Step Functions
#orthogonal polynomial
fit <- lm(wage ~ poly(age, 4), data = Wage)</pre>
coef(summary(fit))
                                                    Pr(>|t|)
                Estimate Std. Error
                                       t value
(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
```

#### raw = TRUE

#it does not afect the ftted values obtained

```
fit2 <- lm(wage ~ poly(age, 4, raw = T), data = Wage)
coef(summary(fit2))
                                         Std. Error
                                                                   Pr(>|t|)
                             Estimate
                                                      t value
(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
#equivalent ways
fit2a \leftarrow lm(wage \sim age + I(age<sup>2</sup>) + I(age<sup>3</sup>) + I(age<sup>4</sup>),
            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)</pre>
age.grid <- seq(from = agelims[1], to = agelims[2]) # #create a grid of values for age
preds <- predict(fit, newdata = list(age = age.grid),</pre>
                  se = TRUE)
se.bands <- cbind(preds$fit + 2 * preds$se.fit,
                   preds$fit - 2 * preds$se.fit)
#plot the data and add the fit from the degree-4 polynomial
par(mfrow = c(1, 2), mar = c(4.5, 4.5, 1, 1),
    oma = c(0, 0, 4, 0)
plot(age, wage, xlim = agelims, cex = .5, col = "darkgrey")
title("Degree -4 Polynomial", outer = T)
lines(age.grid, preds$fit, lwd = 2, col = "blue")
matlines(age.grid, se.bands, lwd = 1, col = "blue", lty = 3)
```

# **Degree -4 Polynomial**



#The ftted values obtained in either case are identical

[1] 1.011813e-11

#ANOVA

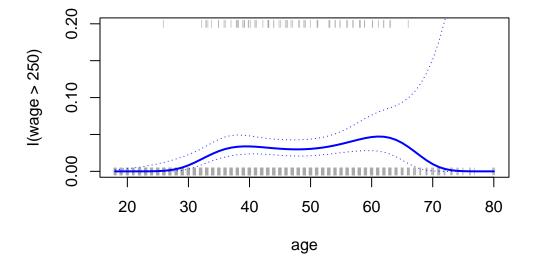
```
fit.1 <- lm(wage ~ age, data = Wage)
fit.2 <- lm(wage ~ poly(age, 2), data = Wage)
fit.3 <- lm(wage ~ poly(age, 3), data = Wage)
fit.4 <- lm(wage ~ poly(age, 4), data = Wage)
fit.5 <- lm(wage ~ poly(age, 5), data = Wage)
anova(fit.1, fit.2, fit.3, fit.4, fit.5)</pre>
```

Analysis of Variance Table

```
Model 1: wage ~ age
Model 2: wage ~ poly(age, 2)
Model 3: wage ~ poly(age, 3)
Model 4: wage ~ poly(age, 4)
Model 5: wage ~ poly(age, 5)
```

```
Res.Df
            RSS Df Sum of Sq F Pr(>F)
   2998 5022216
1
  2997 4793430 1
                      228786 143.5931 < 2.2e-16 ***
2
3 2996 4777674 1
                     15756
                               9.8888 0.001679 **
4 2995 4771604 1
                      6070
                               3.8098 0.051046 .
5 2994 4770322 1
                      1283 0.8050 0.369682
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
coef(summary(fit.5))
               Estimate Std. Error
                                      t value
                                                  Pr(>|t|)
              111.70361 0.7287647 153.2780243 0.000000e+00
(Intercept)
poly(age, 5)1 447.06785 39.9160847 11.2001930 1.491111e-28
poly(age, 5)2 -478.31581 39.9160847 -11.9830341 2.367734e-32
poly(age, 5)3 125.52169 39.9160847 3.1446392 1.679213e-03
poly(age, 5)4 -77.91118 39.9160847 -1.9518743 5.104623e-02
poly(age, 5)5 -35.81289 39.9160847 -0.8972045 3.696820e-01
fit.1 <- lm(wage ~ education + age, data = Wage)</pre>
fit.2 <- lm(wage ~ education + poly(age, 2), data = Wage)</pre>
fit.3 <- lm(wage ~ education + poly(age, 3), data = Wage)</pre>
anova(fit.1, fit.2, fit.3)
Analysis of Variance Table
Model 1: wage ~ education + age
Model 2: wage ~ education + poly(age, 2)
Model 3: wage ~ education + poly(age, 3)
            RSS Df Sum of Sq
 Res.Df
   2994 3867992
1
2 2993 3725395 1
                     142597 114.6969 <2e-16 ***
3 2992 3719809 1
                        5587
                              4.4936 0.0341 *
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
fit \leftarrow glm(I(wage > 250) \sim poly(age, 4), data = Wage,
          family = binomial) #wage > 250 evaluates to a logical variable
```

# directly computed the probabilities by selecting the type = "response" option in the predict() function



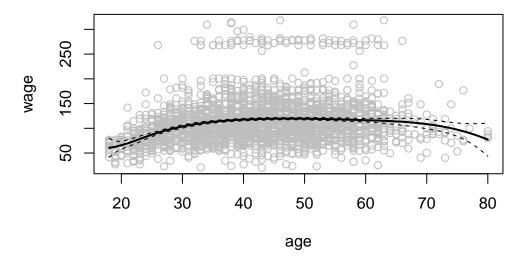
# fit a step function

```
table(cut(age, 4)) #df = 4
(17.9, 33.5]
             (33.5,49]
                        (49,64.5] (64.5,80.1]
        750
                               779
                                            72
                  1399
fit <- lm(wage ~ cut(age, 4), data = Wage)</pre>
coef(summary(fit))
                       Estimate Std. Error t value
                                                         Pr(>|t|)
                      94.158392 1.476069 63.789970 0.000000e+00
(Intercept)
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
```

# Fitting wage to age using a regression spline is simple:

#7.8.2 Splines

```
library(splines)
fit <- lm(wage ~ bs(age, knots = c(25, 40, 60)), data = Wage) #prespecifed knots at ages 25,
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")</pre>
```



```
dim(bs(age, knots = c(25, 40, 60)))
```

[1] 3000 6

```
dim(bs(age, df = 6))
```

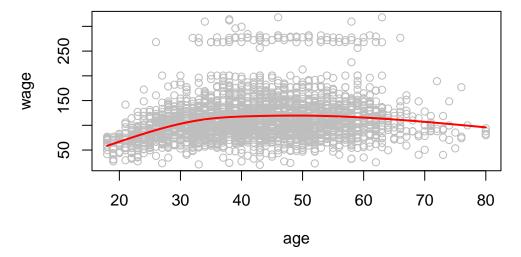
[1] 3000 6

```
attr(bs(age, df = 6), "knots")
```

[1] 33.75 42.00 51.00

#bs() also has a degree argument, so we can ft splines of any degree. default degree = 3 #In order to instead ft a natural spline, we use the ns() function

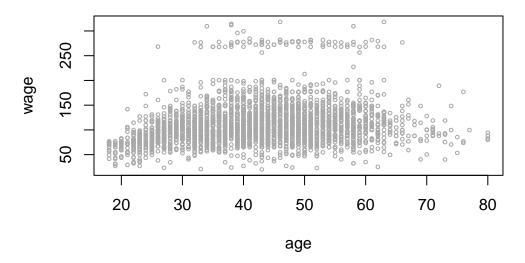
```
fit2 <- lm(wage ~ ns(age, df = 4), data = Wage)
pred2 <- predict(fit2, newdata = list(age = age.grid), se = T)
plot(age, wage, col = "gray")
lines(age.grid, pred2$fit, col = "red", lwd = 2)</pre>
```



#fit a smoothing spline, we use the smooth.spline()

```
plot(age, wage, xlim = agelims, cex = .5, col = "darkgrey")
title("Smoothing Spline")
```

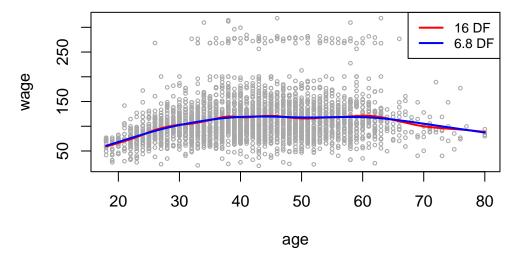
# **Smoothing Spline**



```
fit <- smooth.spline(age, wage, df = 16)
fit2 <- smooth.spline(age, wage, cv = TRUE)</pre>
```

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

#### [1] 6.794596



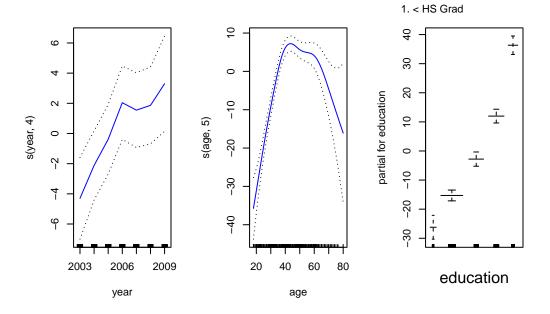
#Notice that in the frst call to smooth.spline(), we specified df = 16. The function then determines which value of leads to 16 degrees of freedom. In the second call to smooth.spline(), we select the smoothness level by crossvalidation; this results in a value of that yields 6.8 degrees of freedom.

#7.8.3 GAMs

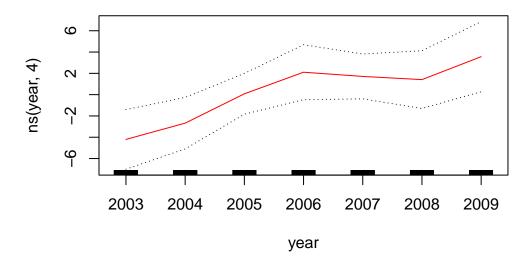
# fit a GAM to predict wage using natural spline functions of year and age

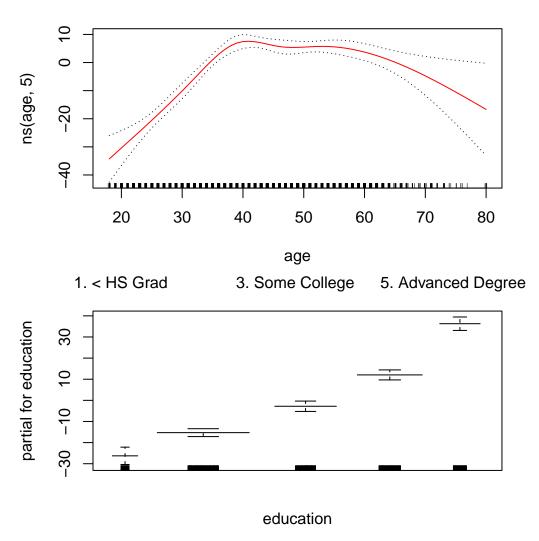
```
gam.m3 <- gam(wage ~ s(year, 4) + s(age, 5) + education, #year should have 4 degrees of free
data = Wage)</pre>
```

```
par(mfrow = c(1, 3))
plot(gam.m3, se = TRUE, col = "blue")
```



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





#:a GAM that excludes year (M1), #a GAM that uses a linear function of year (M2), #a GAM that uses a spline function of year (M3).

#### 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
```

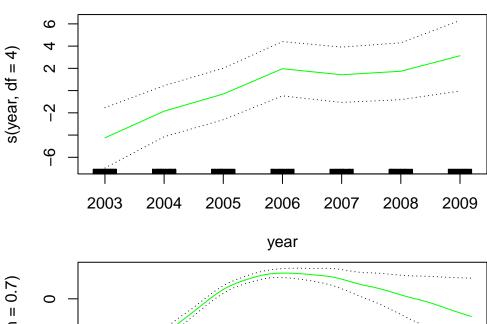
```
Resid. Df Resid. Dev Df Deviance F
                                             Pr(>F)
      2990
              3711731
1
       2989
              3693842 1 17889.2 14.4771 0.0001447 ***
2
3
      2986
              3689770 3 4071.1 1.0982 0.3485661
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
summary(gam.m3)
Call: gam(formula = wage ~ s(year, 4) + s(age, 5) + education, data = Wage)
Deviance Residuals:
            10 Median
    Min
                            3Q
                                   Max
-119.43 -19.70
                -3.33
                         14.17 213.48
(Dispersion Parameter for gaussian family taken to be 1235.69)
   Null Deviance: 5222086 on 2999 degrees of freedom
Residual Deviance: 3689770 on 2986 degrees of freedom
AIC: 29887.75
Number of Local Scoring Iterations: NA
Anova for Parametric Effects
            Df Sum Sq Mean Sq F value
                                          Pr(>F)
                 27162
                        27162 21.981 2.877e-06 ***
s(year, 4)
             1 195338 195338 158.081 < 2.2e-16 ***
s(age, 5)
             4 1069726 267432 216.423 < 2.2e-16 ***
education
Residuals 2986 3689770
                          1236
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Anova for Nonparametric Effects
           Npar Df Npar F Pr(F)
(Intercept)
                 3 1.086 0.3537
s(year, 4)
s(age, 5)
                 4 32.380 <2e-16 ***
education
```

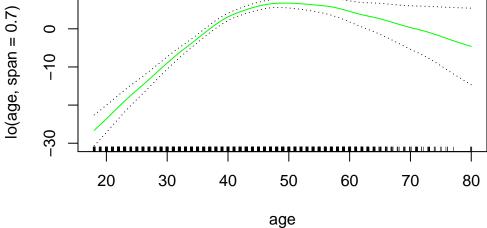
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

```
preds <- predict(gam.m2, newdata = Wage)</pre>
```

#use local regression fts as building blocks in a GAM, using the lo() function #s() function, indicate that we would like to use a smoothing spline

```
gam.lo <- gam(
  wage ~ s(year, df = 4) + lo(age, span = 0.7) + education,
  data = Wage
)
plot(gam.lo, se = TRUE, col = "green")</pre>
```

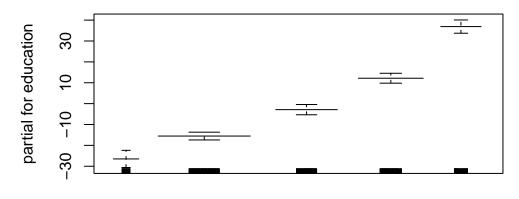




#### 1. < HS Grad

### 3. Some College

## 5. Advanced Degree



#### education

#### library(interp)

#fits a two-term model, in which the first term is an interaction between year and age

Warning in lo.wam(x, z, wz, fit\$smooth, which, fit\$smooth.frame, bf.maxit, : liv too small. (Discovered by lowesd)

Warning in lo.wam(x, z, wz, fit\$smooth, which, fit\$smooth.frame, bf.maxit, : lv too small. (Discovered by lowesd)

Warning in lo.wam(x, z, wz, fit\$smooth, which, fit\$smooth.frame, bf.maxit, : liv too small. (Discovered by lowesd)

Warning in lo.wam(x, z, wz, fit\$smooth, which, fit\$smooth.frame, bf.maxit, : lv too small. (Discovered by lowesd)

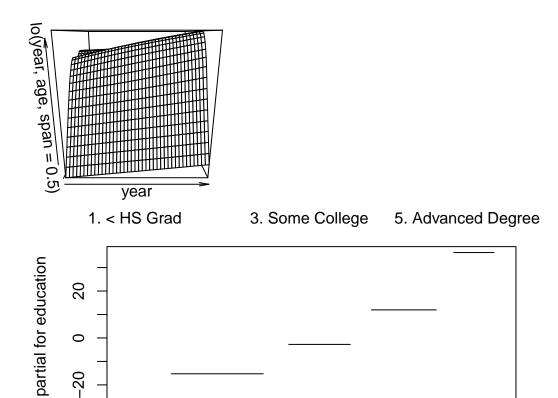
#### library(akima)

Attaching package: 'akima'

```
The following objects are masked from 'package:interp':

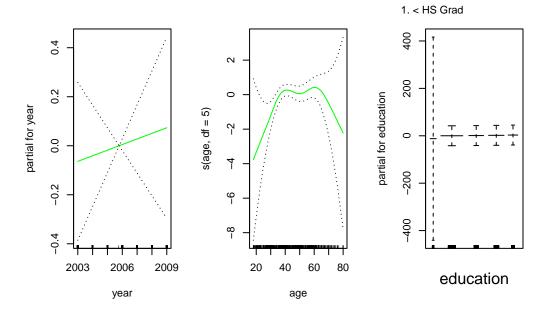
aspline, bicubic, bicubic.grid, bilinear, bilinear.grid,
franke.data, franke.fn, interp, interp2xyz, interpp
```

#### plot(gam.lo.i)



# fit a logistic regression GAM

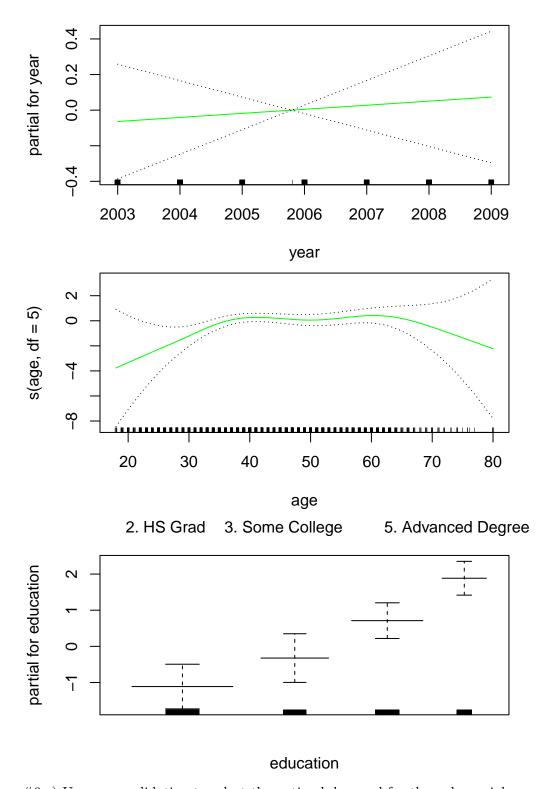
education



#### table(education, I(wage > 250))

education	FALSE	TRUE
1. < HS Grad	268	0
2. HS Grad	966	5
3. Some College	643	7
4. College Grad	663	22
5. Advanced Degree	381	45

#Hence, we ft a logistic regression GAM using all but this category



#6.a) Use cross-validation to select the optimal degree d for the polynomial.

```
set.seed(1)
cv_errors <- rep(NA, 10)

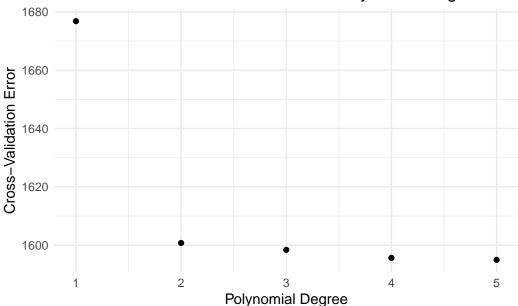
for (d in 1:5) {
    model <- glm(wage ~ poly(age, d), data = Wage)
    cv_errors[d] <- cv.glm(Wage, model, K = 10)$delta[1] #result different from python cause)
}

optimal_degree <- which.min(cv_errors)
print(optimal_degree)</pre>
```

#### [1] 5

Warning: Removed 5 rows containing missing values or values outside the scale range  $(\text{`geom\_point()`})$ .

## Cross-Validation Errors for Different Polynomial Degrees



#6b)

```
cv_errors <- rep(NA, 10)

for (k in 2:12) {
    Wage$age_cut <- cut(Wage$age, k)
    fit <- glm(wage ~ age_cut, data = Wage) # Fit the step function model
    cv_errors[k] <- cv.glm(Wage, fit, K = 10)$delta[1] # Perform 10-fold CV
}

optimal_cuts <- which.min(cv_errors[2:12]) + 1 # +1 because we started at k=2
print(paste("Optimal number of cuts:", optimal_cuts))</pre>
```

[1] "Optimal number of cuts: 11"

```
table(cut(age, 11)) #df = 11
```

```
(17.9,23.6] (23.6,29.3] (29.3,34.9] (34.9,40.5] (40.5,46.2] (46.2,51.8]

143 305 378 503 546 451

(51.8,57.5] (57.5,63.1] (63.1,68.7] (68.7,74.4] (74.4,80.1]

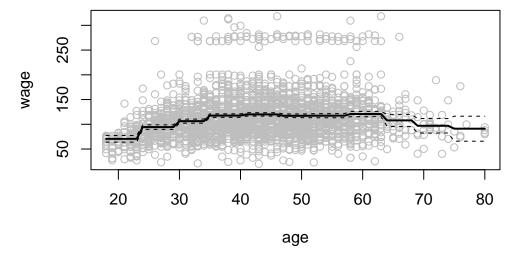
368 223 43 30 10
```

```
fit <- lm(wage ~ cut(age, 11), data = Wage)
coef(summary(fit))</pre>
```

```
Estimate Std. Error
                                              t value
                                                          Pr(>|t|)
(Intercept)
                        70.58039
                                   3.339779 21.133250 1.663885e-92
cut(age, 11)(23.6,29.3] 23.71870
                                             5.859815 5.139182e-09
                                   4.047687
cut(age, 11)(29.3,34.9] 35.76834
                                   3.920945
                                             9.122377 1.315720e-19
cut(age, 11)(34.9,40.5] 47.00149
                                   3.784862 12.418282 1.459360e-34
cut(age, 11)(40.5,46.2] 49.39869
                                   3.751726 13.166924 1.574538e-38
cut(age, 11)(46.2,51.8] 46.57080
                                   3.832858 12.150410 3.415632e-33
cut(age, 11)(51.8,57.5] 46.78093
                                   3.935540 11.886787 7.161447e-32
cut(age, 11)(57.5,63.1] 50.14208
                                   4.278641 11.719159 4.806176e-31
cut(age, 11)(63.1,68.7] 37.29642
                                   6.946084 5.369417 8.504251e-08
cut(age, 11)(68.7,74.4] 26.41431
                                   8.020108
                                             3.293510 1.000974e-03
cut(age, 11)(74.4,80.1] 20.49121
                                  13.063619
                                             1.568571 1.168538e-01
```

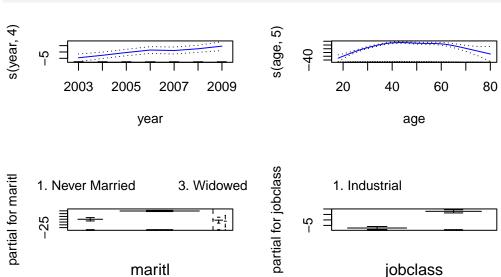
##11 is the same result with python

```
age_grid <- seq(min(Wage$age), max(Wage$age), length.out = 100)
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")</pre>
```



#7. features such as marital status (maritl), job class (jobclass)

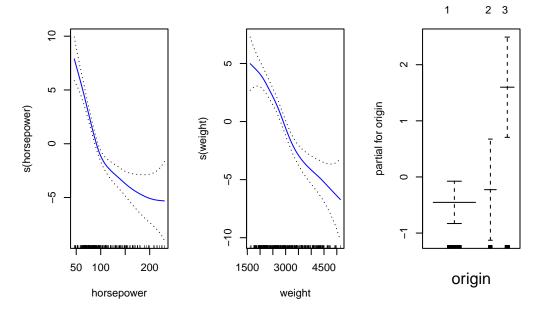
```
par(mfrow = c(2, 2))
plot(gam1, se = TRUE, col = "blue")
```



#same results with python

#8). Fit some of the non-linear models investigated in this chapter to the Auto data set. Is there evidence for non-linear relationships in this data set? Create some informative plots to justify your answer

```
Auto$origin <- factor(Auto$origin)</pre>
```



#9. This question uses the variables dis (the weighted mean of distances to fve Boston employment centers) and nox (nitrogen oxides concentration in parts per 10 million) from the Boston data. We will treat dis as the predictor and nox as the response.

#a)Use the poly() function to fit a cubic polynomial regression to predict nox using dis

```
fit <- lm(nox ~ poly(dis, 3), data = Boston)
summary(fit)</pre>
```

```
Call:
```

lm(formula = nox ~ poly(dis, 3), data = Boston)

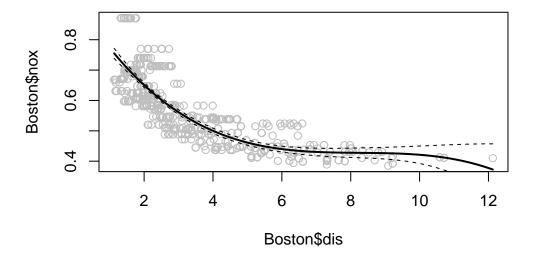
#### Residuals:

Min 1Q Median 3Q Max -0.121130 -0.040619 -0.009738 0.023385 0.194904

#### Coefficients:

```
Estimate Std. Error t value Pr(>|t|)
               0.554695
                          0.002759 201.021
                                             < 2e-16 ***
(Intercept)
poly(dis, 3)1 -2.003096
                          0.062071 -32.271
                                             < 2e-16 ***
poly(dis, 3)2 0.856330
                          0.062071
                                     13.796
                                            < 2e-16 ***
poly(dis, 3)3 -0.318049
                          0.062071
                                     -5.124 4.27e-07 ***
                        0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Signif. codes:
```

```
Residual standard error: 0.06207 on 502 degrees of freedom Multiple R-squared: 0.7148, Adjusted R-squared: 0.7131 F-statistic: 419.3 on 3 and 502 DF, p-value: < 2.2e-16
```



#9b)

```
set.seed(1)

dis_grid <- seq(min(Boston$dis), max(Boston$dis), length.out = 100)

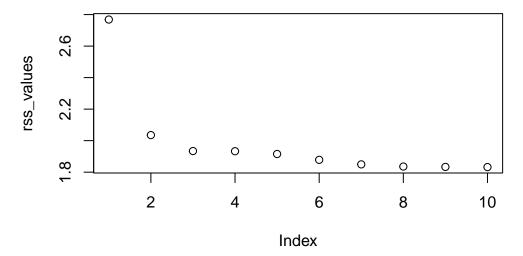
rss_values <- rep(NA, 10)

for (d in 1:10) {
    fit <- glm(nox ~ poly(dis, d), data = Boston)
    preds <- predict(fit, newdata = list(dis = dis_grid))
    rss_values[d] <- sum(residuals(fit)^2)
}

print(rss_values)</pre>
```

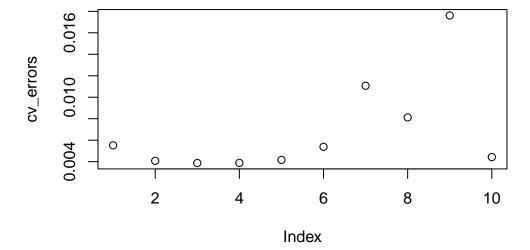
- [1] 2.768563 2.035262 1.934107 1.932981 1.915290 1.878257 1.849484 1.835630
- [9] 1.833331 1.832171

## plot(rss\_values)



```
\#d = 10, 1.832171
\#9c)
```

```
set.seed(1)
cv_errors <- rep(NA, 10)
for (d in 1:10) {
  model <- glm(nox ~ poly(dis, d), data = Boston)
  cv_errors[d] <- cv.glm(Boston, model, K = nrow(Boston))$delta[1]
}
plot(cv_errors)</pre>
```



```
optimal_degree <- which.min(cv_errors)
print(optimal_degree)</pre>
```

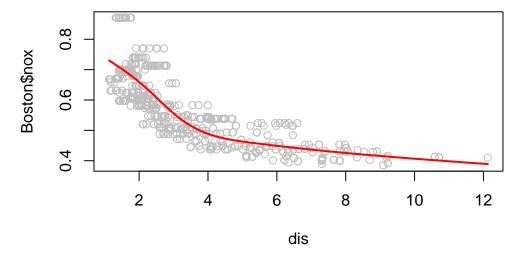
#### [1] 3

#optimal d = 3 with the least cv\_error #9d)

```
attr(bs(Boston$dis, df = 4), "knots")
```

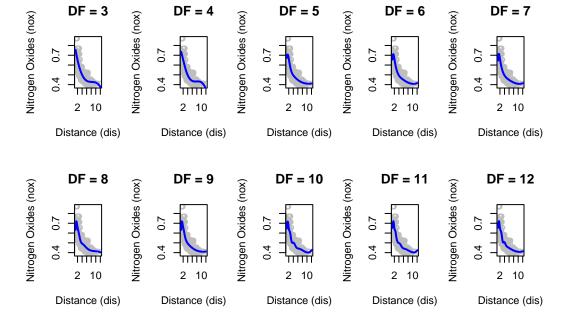
#### [1] 3.20745

```
dis <- Boston$dis
dis.grid <- seq(min(Boston$dis), max(Boston$dis), length.out = 100)
fit <- lm(nox ~ ns(dis, df = 4), data = Boston)
pred <- predict(fit, newdata = list(dis = dis.grid), se = T)
plot(dis, Boston$nox, col = "gray")
lines(dis.grid, pred$fit, col = "red", lwd = 2)</pre>
```



#e)

```
dis_grid <- seq(min(Boston$dis), max(Boston$dis), length.out = 100)
rss_spline <- rep(NA, 10)
par(mfrow = c(2, 5))</pre>
```



```
par(mfrow = c(5, 2))

# Print RSS values for each degree of freedom
print(rss_spline)
```

- [1] 1.934107 1.922775 1.840173 1.833966 1.829884 1.816995 1.825653 1.792535
- [9] 1.796992 1.788999

#9f) Perform cross-validation or another approach in order to select the best degrees of freedom

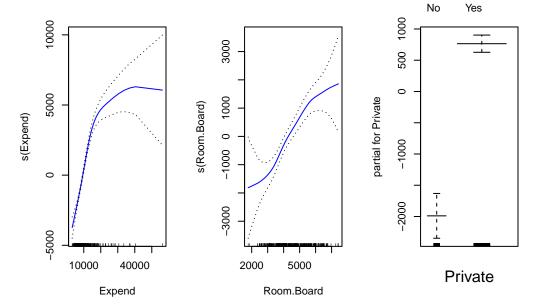
```
#set.seed(1)
#for (df in 3:15) {
  #fit_spline <- glm(nox ~ bs(dis, df = df), data = Boston)</pre>
  #cv_result <- suppressWarnings(cv.glm(Boston, fit_spline))</pre>
  #cv_error_k[df - 2] <- cv_result$delta[1]</pre>
#}
#print("Cross-validation errors (MSE) for different df values (3 to 15):")
#print(cv_error_k)
# Finding the best degrees of freedom based on the lowest MSE
#best_df_index <- which.min(cv_error_k) + 2</pre>
#best_df_index
#10 #a)
y <- College$Outstate
x <- College[, -which(names(College) == "Outstate")]</pre>
set.seed(66)
train <- sample(1:nrow(x), nrow(x) / 2)
test <- (-train)</pre>
y.test <- y[test]</pre>
y.test <- y[test]</pre>
x.train <- x[train, ]</pre>
x.test <- x[test, ]</pre>
y.train <- y[train]</pre>
train_data <- cbind(x.train, Outstate = y.train)</pre>
test_data <- cbind(x.test, Outstate = y.test)</pre>
regfit.fwd <- regsubsets(Outstate ~ ., data = train_data, nvmax = 17, method = "forward")</pre>
summary(regfit.fwd)
Subset selection object
Call: regsubsets.formula(Outstate ~ ., data = train_data, nvmax = 17,
    method = "forward")
17 Variables (and intercept)
             Forced in Forced out
```

```
PrivateYes
                   FALSE
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                   FALSE
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Accept
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Enroll
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Top10perc
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Top25perc
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P.Undergrad
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```

#Room. Board #expend #PrivateYes the predictors change if the training sample change #but the 2/3 of the predictors are the same with python

#(b) Fit a GAM on the training data, using out-of-state tuition as the response and the features selected in the previous step as the predictors. Plot the results, and explain your findings



#11.

```
# a)
set.seed(66)
X1 <- rnorm(100, mean = 0, sd = 1)
X2 <- rnorm(100, mean = 0, sd = 1)
Y <- 3 + 2 * X1 + 3 * X2 + rnorm(100, mean = 0, sd = 1)</pre>
```

```
#b)
beta1 <- 0

#c)
a <- Y - beta1 * X1
beta2 <- lm(a ~ X2)$coef[2]

#d)
a <- Y - beta2 * X2
beta1 <- lm(a ~ X1)$coef[2]

#e)
iterations <- 600
beta0_vals <- beta1_vals <- beta2_vals <- numeric(iterations)
beta2 <- 0</pre>
```

```
for (i in 1:iterations) {
    # (c) Keep beta1 fixed, fit Y - beta1*X1 = beta0 + beta2*X2 + error
    a <- Y - beta1 * X1
    beta2 <- lm(a ~ X2)$coef[2] # Update beta2

# (d) Keep beta2 fixed, fit Y - beta2*X2 = beta0 + beta1*X1 + error
    a <- Y - beta2 * X2
    beta1 <- lm(a ~ X1)$coef[2] # Update beta1

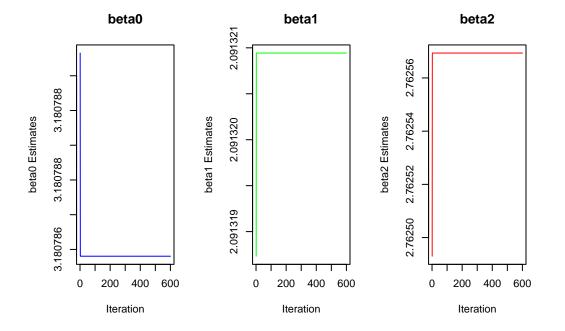
# Store beta estimates for each iteration
    beta0_vals[i] <- lm(a ~ X1)$coef[1]
    beta1_vals[i] <- beta1
    beta2_vals[i] <- beta2
}</pre>
```

```
par(mfrow = c(1, 3))

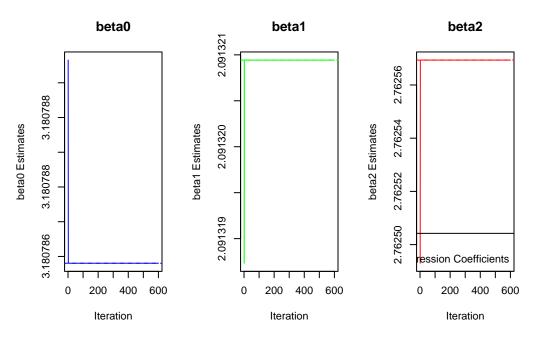
plot(1:iterations, beta0_vals, type = "l", col = "blue",
        ylab = "beta0 Estimates", xlab = "Iteration", main = "beta0")

plot(1:iterations, beta1_vals, type = "l", col = "green",
        ylab = "beta1 Estimates", xlab = "Iteration", main = "beta1")

plot(1:iterations, beta2_vals, type = "l", col = "red",
        ylab = "beta2 Estimates", xlab = "Iteration", main = "beta2")
```



```
# f)
fit <-lm(Y \sim X1 + X2)
summary(fit)
Call:
lm(formula = Y \sim X1 + X2)
Residuals:
    Min
            1Q Median
                            3Q
                                   Max
-3.6423 -0.7321 -0.0059 0.8477 3.1870
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept)
             3.1808
                      0.1058 30.06
                                         <2e-16 ***
Х1
             2.0913
                        0.1094 19.12 <2e-16 ***
X2
             2.7626
                        0.1211 22.81
                                         <2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
Residual standard error: 1.056 on 97 degrees of freedom
Multiple R-squared: 0.8985, Adjusted R-squared: 0.8965
F-statistic: 429.5 on 2 and 97 DF, p-value: < 2.2e-16
par(mfrow = c(1, 3))
plot(1:iterations, beta0_vals, type = "l", col = "blue",
     ylab = "beta0 Estimates", xlab = "Iteration", main = "beta0")
abline(h = coef(fit)[1], col = "blue", lty = 2)
plot(1:iterations, beta1_vals, type = "1", col = "green",
     ylab = "beta1 Estimates", xlab = "Iteration", main = "beta1")
abline(h = coef(fit)[2], col = "green", lty = 2)
plot(1:iterations, beta2_vals, type = "l", col = "red",
     ylab = "beta2 Estimates", xlab = "Iteration", main = "beta2")
abline(h = coef(fit)[3], col = "red", lty = 2)
legend("bottomright", legend = c("Backfitting", "Multiple Regression Coefficients"),
       col = c("black", "black"), lty = c(1, 2))
```



```
tolerance <- 0.00001

fit <- lm(Y ~ X1 + X2)
beta0_mlr <- coef(fit)[1]
beta1_mlr <- coef(fit)[2]
beta2_mlr <- coef(fit)[3]

# Check convergence in each iteration
for (i in 1:iterations) {
   if (abs(beta0_vals[i] - beta0_mlr) < tolerance &&
        abs(beta1_vals[i] - beta1_mlr) < tolerance &&
        abs(beta2_vals[i] - beta2_mlr) < tolerance) {
        cat("Convergence reached at iteration", i, "\n")
        break
   }
}</pre>
```

### Convergence reached at iteration 2

```
if (i == iterations) {
  cat("No convergence within the tolerance of", tolerance, "after", iterations, "iterations."
}
```