# STATS 415 Lab9

Weijing Tang Mar 15, 2018

## 1 Today's objectives

- 1. Learn how to implement polynomial regression.
- 2. Learn how to implement regression spline and smoothing spline.
- 3. Learn how to implement GAM.
- 4. Compare the model performance among polynomial regression, splines and GAM.

## 2 Non-Linear Methods.

So far we have focused on linear models. They have a significant advantage: simple. It is great for interpretation and inference. In this lab, we will implement several non-linear methods whose goal is to relax the linearity assumption but attempt to maintain as much interpretability as possible.

We will use the Wage dataset from the ISLR package to explore the following three methods:

- Polynomial regression
- Splines
- Generalized additive models(GAM)

This dataset contains wage and other data for a group of 3000 male workers. We want to predict wage here.

```
library(ISLR)
attach(Wage)
```

# 3 Spliting training and test data

We will just use training data to select the relevent "degree of freedom" parameter for each model, then compare their performance on test data.

```
set.seed(1)
train = sample(1:nrow(Wage),trunc(nrow(Wage)*0.8))
```

# 4 Polynomial Regression

We extent linear regression to

$$y = \beta_0 + \beta_1 x + \beta_2 x^2 + \dots + \beta_d x^d + \epsilon$$

where d decides the degree of freedom of the model.

We will use age to predict wage. In order to predict wage using a d-degree polynomial in age, we need create a matrix which contains  $[x, x^1, \dots, x^d]$  for each observation. Then we can use lm() function to fit a linear model as what we learned in the previous labs. poly() returns a matrix whose columns are a basis of orthogonal polynomials, which essentially means that each column is a linear orthogonal combination of the variables  $[x, x^1, \dots, x^d]$ . It is more numerically stable compared with the original matrix.

However, we can also use poly() to obtain [age,age^2,age^3,age^4] directly by adding raw=TRUE argument. Then the estimated coefficients has a clear interpretation. Later we will show that this change does not affect the fitted model (give the same prediction).

2.787932 5.346449e-03

```
fit2=lm(wage~poly(age,4,raw=T),data=Wage[train,])
coef(summary(fit2))
```

## poly(age, 4)4 -71.15018 40.2724169 -1.766722 7.740207e-02

It is equivalent to the following model:

## poly(age, 4)3 112.27676 40.2724169

```
fit2a=lm(wage~age+I(age^2)+I(age^3)+I(age^4),data=Wage[train,])
coef(fit2a)
```

```
## (Intercept) age I(age^2) I(age^3) I(age^4)
## -1.891721e+02 2.157842e+01 -5.692226e-01 6.837459e-03 -3.219978e-05
```

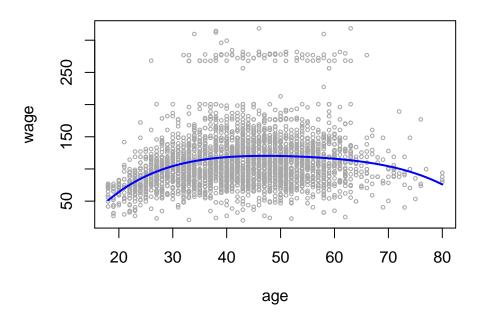
You can see the coefficient estimators are the same.

#### 4.1 visualization

To visualize the fitted curve, we create a grid of values for age at which we make predictions by using generic predict() function.

```
agelims=range(age)
age.grid=seq(from=agelims[1],to=agelims[2])
preds=predict(fit,newdata=data.frame(age=age.grid))
plot(age,wage,xlim=agelims,cex=.5,col="darkgrey")
title("Degree-4 Polynomial")
lines(age.grid,preds,lwd=2,col="blue")
```

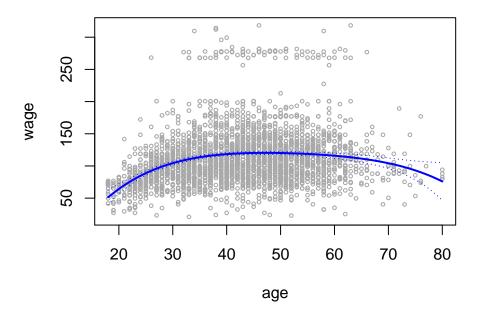
# **Degree-4 Polynomial**



We can also include standard errors in the plot by using se=TRUE argument of predict() function. Here we choose 95% confidence interval, so the band should be within roughly 1.96 standard errors of predicted values.

```
preds=predict(fit,newdata=data.frame(age=age.grid),se=TRUE)
se.bands=cbind(preds$fit+1.96*preds$se.fit,preds$fit-1.96*preds$se.fit)
plot(age,wage,xlim=agelims,cex=.5,col="darkgrey")
title("Degree-4 Polynomial")
lines(age.grid,preds$fit,lwd=2,col="blue")
lines(age.grid,se.bands[,1],lwd=1,col="blue",lty=3)
lines(age.grid,se.bands[,2],lwd=1,col="blue",lty=3)
```

## **Degree-4 Polynomial**



We mentioned earlier that whether or not an orthogonal set of basis functions is produced in the poly() function will not affect the fitted model. It means the fitted values obtained in either case are identical:

```
preds2=predict(fit2,newdata=data.frame(age=age.grid),se=TRUE)
max(abs(preds$fit-preds2$fit))
```

## [1] 5.151435e-11

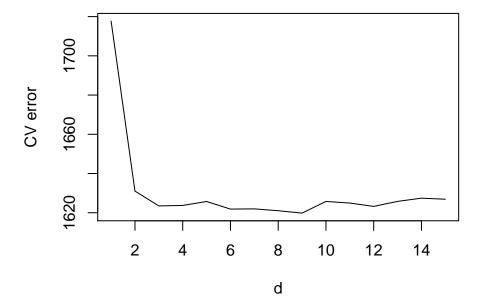
#### 4.2 How to select the degree d?

In performing a polynomial regression we must decide on the degree of the polynomial to use. We will implement two methods. One way is to choose d by cross-validation. We have practiced it in Lab6. Let's go over it again!

We will perform linear regression using the glm() function rather than the lm() function because the latter can be used together with cv.glm(). cv.glm() function is in package boot. We usually use 10-fold cross-validation by setting K=10 argument. The standard cross-validation error is saved as delta[1] component in the returned list.

```
library(boot)
set.seed(1)
cv.error_poly = rep(0,15)
for (i in 1:15){
   fit=glm(wage~poly(age,i),data=Wage[train,])
    cv.error_poly[i]=cv.glm(Wage[train,], fit, K=10)$delta[1]
}
cv.error_poly
```

```
## [1] 1717.707 1631.020 1623.470 1623.707 1625.740 1621.858 1621.958
## [8] 1621.026 <mark>1619.813</mark> 1625.750 1624.945 1623.216 1625.782 1627.430
```



Question: Which degree will you choose based on the result of cross-validation?

Another way is to choose d by using hypothesis tests.

$$y = \beta_0 + \beta_1 x + \beta_2 x^2 + \dots + \beta_d x^d + \epsilon$$

For example, the null hypothesis is  $H_0: \beta_{k+1} = \cdots = \beta_d = 0$ . If we reject the null hypothesis, we think the polynomial regression with degree k is sufficient to explain the data against a more complex model. In general we use anova() function to test the null hypothesis that a model  $M_1$  is sufficient to explain the data against the alternative hypothesis that a more complex model  $M_2$  is required.  $M_1$  and  $M_2$  should be nested models if we use anova() function, which means that the predictors in  $M_1$  must be a subset of the predictors in  $M_2$ . In this case, we fit models ranging from linear to a degree-5 polynomial and sequentially compare the simpler model to the more complex model, determine the simplest model which is sufficient to explain the relationship between wage and age.

```
fit.1=lm(wage~age,data=Wage[train,])
fit.2=lm(wage~poly(age,2),data=Wage[train,])
fit.3=lm(wage~poly(age,3),data=Wage[train,])
fit.4=lm(wage~poly(age,4),data=Wage[train,])
fit.5=lm(wage~poly(age,5),data=Wage[train,])
anova(fit.1,fit.2,fit.3,fit.4,fit.5)
```

```
## 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
                                            Pr(>F)
## 1
      2398 4117772
## 2
      2397 3902041 1
                         215730 133.0405 < 2.2e-16 ***
      2396 3889435 1
                          12606
                                  7.7741 0.005342 **
                           5062
                                  3.1219 0.077372 .
## 4
      2395 3884373 1
      2394 3881967 1
                           2406
                                  1.4838 0.223294
## 5
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

Question: Which degree will you choose based on the result of anova(or p-values)? Which degree will you choose finally?

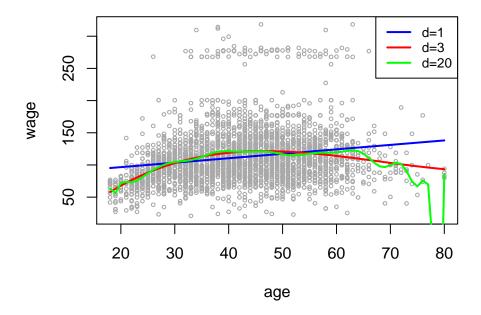
We prefer to choose 3. Because after 3 the error is quite similar and the model for 3 is simpler

### 4.3 What kind of role does degree d play in the model?

Let's visualize the fitted curve with different degrees in one plot!

```
fit.20=lm(wage~poly(age,20),data=Wage[train,])
preds1=predict(fit.1,newdata=data.frame(age=age.grid))
preds3=predict(fit.3,newdata=data.frame(age=age.grid))
preds20=predict(fit.20,newdata=data.frame(age=age.grid))
plot(age,wage,xlim=agelims,cex=.5,col="darkgrey")
title("Polynomial regression")
lines(age.grid,preds1,lwd=2,col="blue")
lines(age.grid,preds3,lwd=2,col="red")
lines(age.grid,preds20,lwd=2,col="green")
legend("topright",legend=c("d=1","d=3","d=20"),col=c("blue","red","green"),lty=1,lwd=2,cex=.8)
```

# **Polynomial regression**



Question: How does the fitted curve change as we increase the degree d?

overfit

# 5 Splines

The key idea of regression splines is to fit different polynomials locally (over different regions of x). Fortunately we can represent a spline with k knots by constructing an appropriate matrix of basis functions.

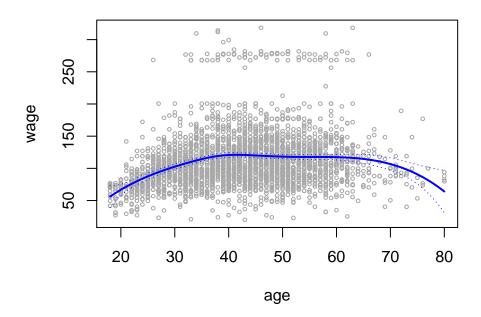
### 5.1 Cubic splines

The bs() function generates the entire matrix for splines with the specified set of knots, which is in package splines. The degree of freedom is decided by the degree of polynomial and the number of knots. We can set degree=d and knots=... or set df= directly. By default, degree=3 for cubic splines. In the case of cubic spline, df = #knots + 4. In practice, knots are often placed at uniform quantiles of the data. For example, if we just set df=7, bs() function will generate the matrix for splines with knots placed at 0.25, 0.5, 0.75 quantiles of age. This produces a spline with 7 basis functions.(including the intercept term)

```
library(splines)
fit=lm(wage~bs(age,df = 7),data=Wage[train,])
```

```
preds=predict(fit,newdata=data.frame(age=age.grid),se=TRUE)
se.bands=cbind(preds$fit+1.96*preds$se.fit,preds$fit-1.96*preds$se.fit)
plot(age,wage,xlim=agelims,cex=.5,col="darkgrey")
title("Cubic Spline with df=7")
lines(age.grid,preds$fit,lwd=2,col="blue")
lines(age.grid,se.bands[,1],lwd=1,col="blue",lty=3)
lines(age.grid,se.bands[,2],lwd=1,col="blue",lty=3)
```

# **Cubic Spline with df=7**



We can also use specified knots.

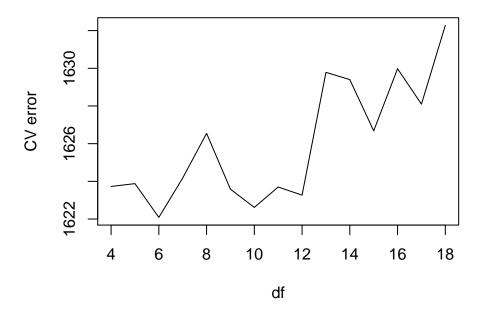
```
dim(bs(age,knots=c(25,40,60)))
## [1] 3000 6
```

### 5.2 How to choose the degree of freedom of splines?

```
We can choose df by cross-validation. degree!=df
```

```
set.seed(1)
cv.error_cs = rep(0,15)
for (i in 1:15){
  fit=glm(wage~bs(age,df = i+3),data=Wage[train,])
  cv.error_cs[i]=cv.glm(Wage[train,], fit, K=10)$delta[1]
}
which.min(cv.error_cs)
```

```
## [1] 3
plot(4:18,cv.error_cs,xlab = "df",ylab = "CV error",type = "l")
```



```
fit.cs = glm(wage~bs(age,df = 6),data=Wage[train,])
```

Question: How many knots will you choose based on the result of cross-validation?

### 5.3 Natural splines

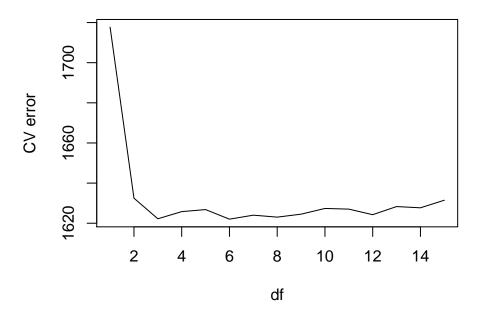
Natural spline replaces the "end" cubic splines (one on each side) with lines which gains stability on the boundary. Refer to the plot on Page 24 in the lecture notes. We use ns() function instead of bs(). ns() function works similarly as bs(). In the case of natural spline, df = #knots + 1.

Let's choose the degree of freedom again by cross-validation.

```
set.seed(1)
cv.error_ns = rep(0,15)
for (i in 1:15){
  fit=glm(wage~ns(age,df = i),data=Wage[train,])
  cv.error_ns[i]=cv.glm(Wage[train,], fit, K=10)$delta[1]
}
which.min(cv.error_ns)
```

## [1] 6

plot(1:15,cv.error\_ns,xlab = "df",ylab = "CV error",type = "l")



fit.ns = glm(wage~ns(age,df = 6),data=Wage[train,])

### 5.4 Smoothing spline

The key idea of smooth splines is to find some function g that makes RSS small but is also smooth. The objective function is

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

where the smoothness of g is controlled by tuning parameter  $\lambda$ .

**Question:** What kind of model we are fitting if  $\lambda = \infty$ ?

We don't give a explicit form of the degree of freedom, but it is uniquely defined by  $\lambda$ . In general, the larger  $\lambda$  is, the smaller the degree of freedom is. We use the smooth.spline() function to fit a smoothing spline.

fit=smooth.spline(age,wage,df=16)

We can do cross-validation and visualization by using smooth.spline() function directly.

```
fit.ss=smooth.spline(age,wage,cv=TRUE)

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

fit.ss$df

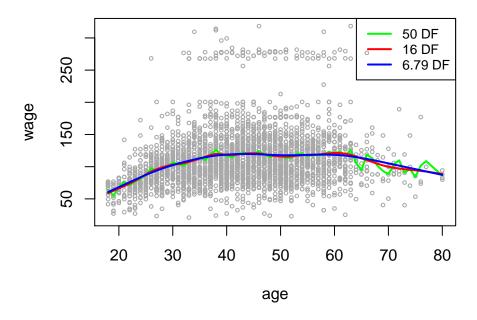
## [1] 6.794596

fit.ss$lambda

## [1] 0.02792303

fit.50=smooth.spline(age,wage,df=50)
plot(age,wage,xlim=agelims,cex=.5,col="darkgrey")
title("Smoothing Spline")
lines(fit.50,col="green",lwd=2)
lines(fit,col="red",lwd=2)
lines(fit.ss,col="blue",lwd=2)
legend("topright",legend=c("50 DF","16 DF","6.79 DF"),col=c("green","red","blue"),lty=1,lwd=2,cex=.8)
```

# **Smoothing Spline**



Question: How does the fitted curve change as we increase the degree of freedom?

## 6 GAMs

GAM replaces each term in the multiple linear regression model with a non-linear function:

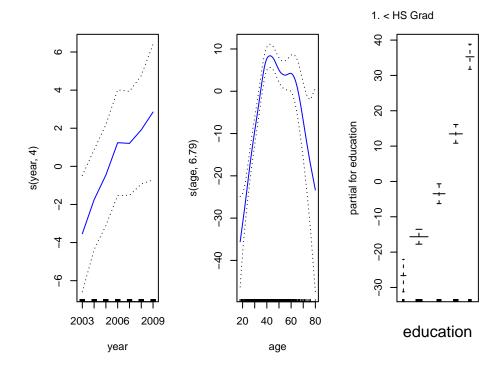
$$y = \beta_0 + f_1(x_1) + f_2(x_2) + \dots + f_p(x_p) + \epsilon$$

We now fit a GAM to predict wage using natural spline functions of year and age, treating education as a qualitative predictor. We can create spline matrix of basis functions by using ns(), bs() functions and use lm() function.

```
gam1=lm(wage~ns(year,4)+ns(age,5)+education,data=Wage[train,])
```

In order to fit more general sorts of GAMs, we will need to use the gam library in R. The s() function, which is part of the gam library, is used to indicate that we would like to use a smoothing spline. The second argument is the target degree of freedom. Here we use smoothing spline with df = 6.79 for predictor age. Let's also try smoothing spline for year.

```
library(gam)
gam2=gam(wage~s(year,4)+s(age,6.79)+education,data=Wage[train,])
par(mfrow=c(1,3))
plot(gam2, se=TRUE,col="blue")
```



Notice that if we want to plot a GAM fitted object obtained from lm(), we had to use plot.gam() rather than the generic plot() function.

In these plots, the function of year looks rather linear. We can perform ANOVA test in order to determine which of the models is better: a GAM that uses a linear function of year  $(M_1)$ , or a GAM that uses a spline function of year  $(M_2)$ .

```
gam.m1=gam(wage~year+s(age,6.79)+education,data=Wage[train,])
gam.m2=gam(wage~s(year,4)+s(age,6.79)+education,data=Wage[train,])
anova(gam.m1,gam.m2,test="F")
```

```
## Analysis of Deviance Table
##
## Model 1: wage ~ year + s(age, 6.79) + education
## Model 2: wage ~ s(year, 4) + s(age, 6.79) + education
## Resid. Df Resid. Dev Df Deviance F Pr(>F)
## 1 2387.2 3004854
## 2 2384.2 3003602 3 1251.4 0.3311 0.8029
```

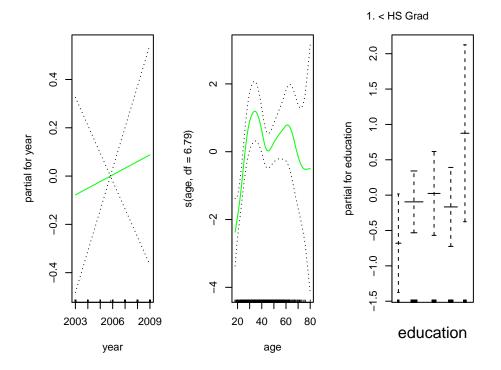
Question: Based on the p-value in the anova result, which model is preferred?

We can make predictions from gam objects, just like from 1m objects, using the predict() method for the class gam.

## 6.1 logistic GAM

We set argument family=binomial as what we did for logistic regression using glm() function. In order to fit a logistic regression GAM, we use the I() function in constructing the binary response variable.

```
gam.lr=gam(I(wage>50) ~ year + s(age,df=6.79) + education, family=binomial,data=Wage[train,])
par(mfrow=c(1,3))
plot(gam.lr,se=T,col="green")
```



7 Compare the performance of polynomial regression, cubic spline, natural spline and smoothing spline

```
test.poly = predict(fit.poly,Wage[-train,])
test.cs = predict(fit.cs,Wage[-train,])
test.ns = predict(fit.ns,Wage[-train,])
test.ss = predict(fit.ss,age[-train])
test.gam = predict(gam.m1,Wage[-train,])
wage.test = wage[-train]
error.poly = mean(wage.test-test.poly)
error.cs = mean(wage.test-test.cs)
error.ns = mean(wage.test-test.ns)
error.ss = mean(wage.test-test.ss$y)
error.gam = mean(wage.test-test.gam)
d <- data.frame("TestMSE" = c(error.poly, error.cs, error.ns, error.ss, error.gam))
rownames(d) <- c("poly regression", "cubic spline", "natural spline", "smoothing spline", "GAM")
knitr::kable(d)</pre>
```

	TestMSE
poly regression	-2.275973
cubic spline	-2.392264
natural spline	-2.406441
smoothing spline	-1.893247
GAM	-1.434419

Question: Which model do you prefer from the table above?

### 8 Resources and References

1. Partially adapted from http://www-bcf.usc.edu/~gareth/ISL/Chapter%207%20Lab.txt