

Solutions to Assignment

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Conceptual 1

(a)

$$\begin{aligned} f(x) &= f_1(x) \\ \beta_0 + \beta_1 x + \beta_2 x^2 + \beta_3 x^3 + \beta_4 (x - \xi)_+^3 &= a_1 + b_1 x + c_1 x^2 + d_1 x^3 \\ (\beta_0 - a_1) + (\beta_1 - b_1)x + (\beta_2 - c_1)x^2 + (\beta_3 - d_1)x^3 &= 0 \end{aligned} \quad (x - \xi)_+^3 = 0$$

$$\beta_0 = a_1$$

$$\beta_1 = b_1$$

$$\beta_2 = c_1$$

$$\beta_3 = d_1$$

(b)

$$\begin{aligned} f(x) &= f_2(x) \\ \beta_0 + \beta_1 x + \beta_2 x^2 + \beta_3 x^3 + \beta_4 (x - \xi)_+^3 &= a_2 + b_2 x + c_2 x^2 + d_2 x^3 \\ \beta_0 + \beta_1 x + \beta_2 x^2 + \beta_3 x^3 + \beta_4 (x^3 - 3\xi x^2 + 3\xi^2 x - \xi^3) &= a_2 + b_2 x + c_2 x^2 + d_2 x^3 \\ (\beta_0 - a_2 - \beta_4 \xi^3) + (\beta_1 - b_2 + 3\beta_4 \xi^2)x + (\beta_2 - c_2 - 3\beta_4 \xi)x^2 + (\beta_3 - d_2 + \beta_4)x^3 &= 0 \end{aligned}$$

$$a_2 = \beta_0 - \beta_4 \xi^3$$

$$b_2 = \beta_1 + 3\beta_4 \xi^2$$

$$c_2 = \beta_2 - 3\beta_4 \xi$$

$$d_2 = \beta_3 + \beta_4$$

(c)

$$\begin{aligned} f_2(\xi) &= \beta_0 - \beta_4 \xi^3 + (\beta_1 + 3\beta_4 \xi^2)\xi + (\beta_2 - 3\beta_4 \xi)\xi^2 + (\beta_3 + \beta_4)\xi^3 \\ &= \beta_0 + \beta_1 \xi + \beta_2 \xi^2 + \beta_3 \xi^3 \\ &= f_1(\xi) \end{aligned}$$

(d)

$$\begin{aligned} f_1'(\xi) &= \beta_1 + 2\beta_2 \xi + 3\beta_3 \xi^2 \\ f_2'(\xi) &= \beta_1 + 3\beta_4 \xi^2 + 2(\beta_2 - 3\beta_4 \xi)\xi + 3(\beta_3 + \beta_4)\xi^2 \\ &= \beta_1 + 2\beta_2 \xi + 3\beta_3 \xi^2 \\ &= f_1'(\xi) \end{aligned}$$

(e)

$$f_1''(x) = 2\beta_2 + 6\beta_3x$$

$$\begin{aligned} f_2''(x) &= 2\beta_2 - 6\beta_4\xi + 6(\beta_3 + \beta_4)\xi^2 \\ &= f_1''(x) \end{aligned}$$

Conceptual 2

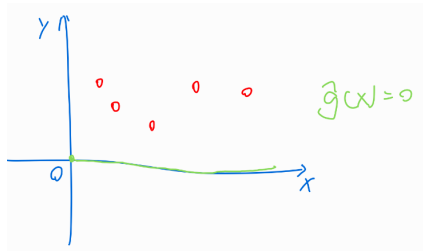


Figure 1: (a)

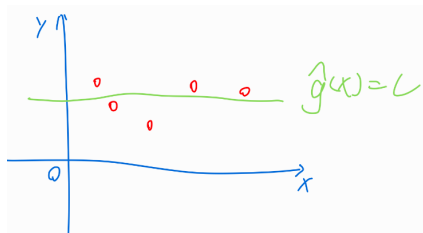


Figure 2: (b)

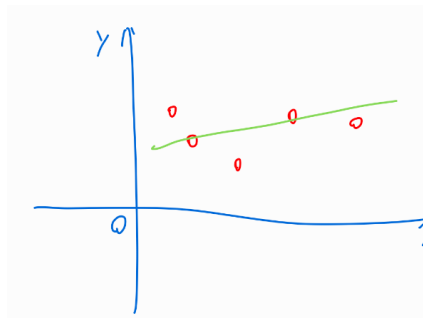


Figure 3: (c)

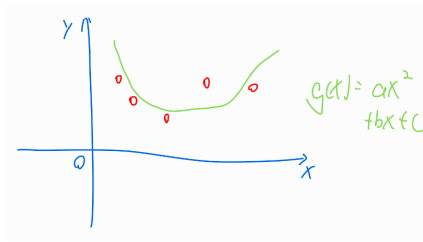


Figure 4: (d)

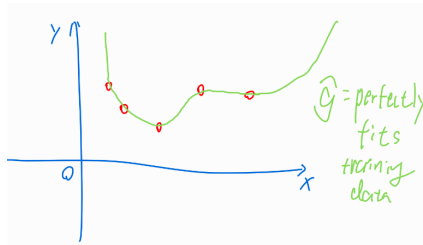


Figure 5: (e)

Conceptual 3

$$1/x \quad \text{and} \quad 1/x - 2(x-2)^2$$

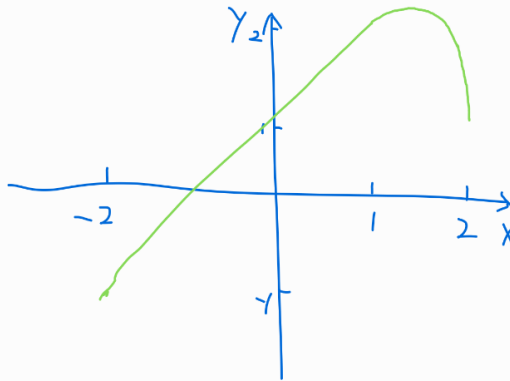


Figure 6: 3

Conceptual 5

- (a) g_2 because it is more flexible and can fit the data better.
- (b) cannot be determined. However, if g_2 overfits then g_1 will perform better.
- (c) same since the penalty term

Applied 6

a

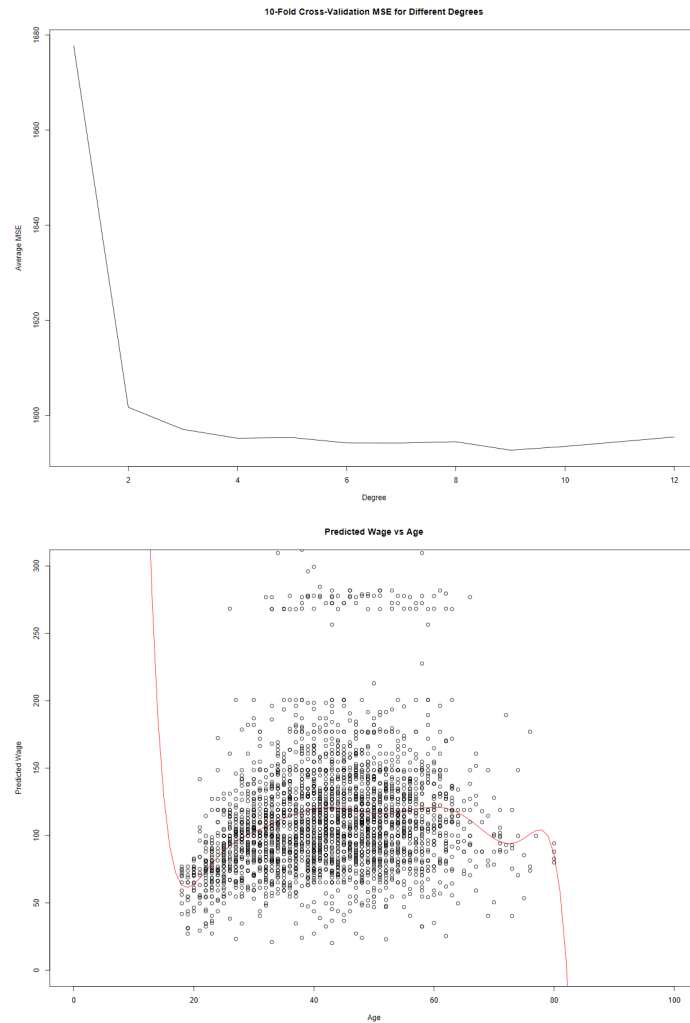


Figure 7

```
library(ISLR2)

data(Wage)
min(Wage$age)
max(Wage$age)

cv.poly <- function(k = 10, Wage, degs) {
  # get fixed indices for folds and shuffle
  indices <- c(1:nrow(Wage))
  indices <- sample(indices)

  # split into k folds
  folds <- split(indices, rep(1:k, length.out = length(indices)))

  avg_mse <- c()
  for (deg in degs) {
```

```

mse <- c()
for (fold in folds) {
  fit <- lm(wage ~ poly(age, degree = deg), data = Wage[-fold, ])
  pred <- predict(fit, Wage[fold, ])
  mse <- c(mse, mean((pred - Wage[fold, ]$wage)^2))
}
avg_mse <- c(avg_mse, mean(mse))
}

# find the best index
best_index <- which.min(avg_mse)
return(list(best_deg = degs[best_index], avg_mse = avg_mse, degs = degs))
}

cv_results <- cv.poly(k = 10, Wage, degs = 2:12)
cv_results$best_deg
# best degree is 9

plot(cv_results$avg_mse ~ cv_results$degs, type = "l", xlab = "Degree", ylab = "Average MSE", main = "10-Fold")

# best model
best_model <- lm(wage ~ poly(age, degree = 9), data = Wage)
summary(best_model)

pred <- predict(best_model, data.frame(age = seq(0, 100, 1)))

# xlim from 0 to 100
# ylim from 0 to 300
plot(pred ~ seq(0, 100, 1), xlab = "Age", ylab = "Predicted Wage", main = "Predicted Wage vs Age", type = "l")
points(Wage$age, Wage$wage)

models <- lapply(1:12, function(deg) lm(wage ~ poly(age, degree = deg), data = Wage))
# anova all models
do.call(anova, models)

```

Coefficients:				
	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	111.7036	0.7282	153.395	< 2e-16 ***
poly(age, degree = 9)1	447.0679	39.8857	11.209	< 2e-16 ***
poly(age, degree = 9)2	-478.3158	39.8857	-11.992	< 2e-16 ***
poly(age, degree = 9)3	125.5217	39.8857	3.147	0.00167 **
poly(age, degree = 9)4	-77.9112	39.8857	-1.953	0.05087 .
poly(age, degree = 9)5	-35.8129	39.8857	-0.898	0.36932
poly(age, degree = 9)6	62.7077	39.8857	1.572	0.11601
poly(age, degree = 9)7	50.5498	39.8857	1.267	0.20512
poly(age, degree = 9)8	-11.2547	39.8857	-0.282	0.77783
poly(age, degree = 9)9	-83.6918	39.8857	-2.098	0.03596 *

Figure 8

	Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)	
1	2998	5022216					
2	2997	4793430	1	228786	143.6709	< 2.2e-16	***
3	2996	4777674	1	15756	9.8941	0.001674	**
4	2995	4771604	1	6070	3.8119	0.050984	.
5	2994	4770322	1	1283	0.8054	0.369552	
6	2993	4766389	1	3932	2.4693	0.116192	
7	2992	4763834	1	2555	1.6046	0.205345	
8	2991	4763707	1	127	0.0795	0.777935	
9	2990	4756703	1	7004	4.3985	0.036054	*
10	2989	4756701	1	3	0.0017	0.967540	
11	2988	4756597	1	103	0.0648	0.799070	
12	2987	4756591	1	7	0.0043	0.947903	

Figure 9

the best degree is 9. The anova tests show that only degree 2 3 and 9 significantly improve the degree 1 model. This is consistent with summary output

b

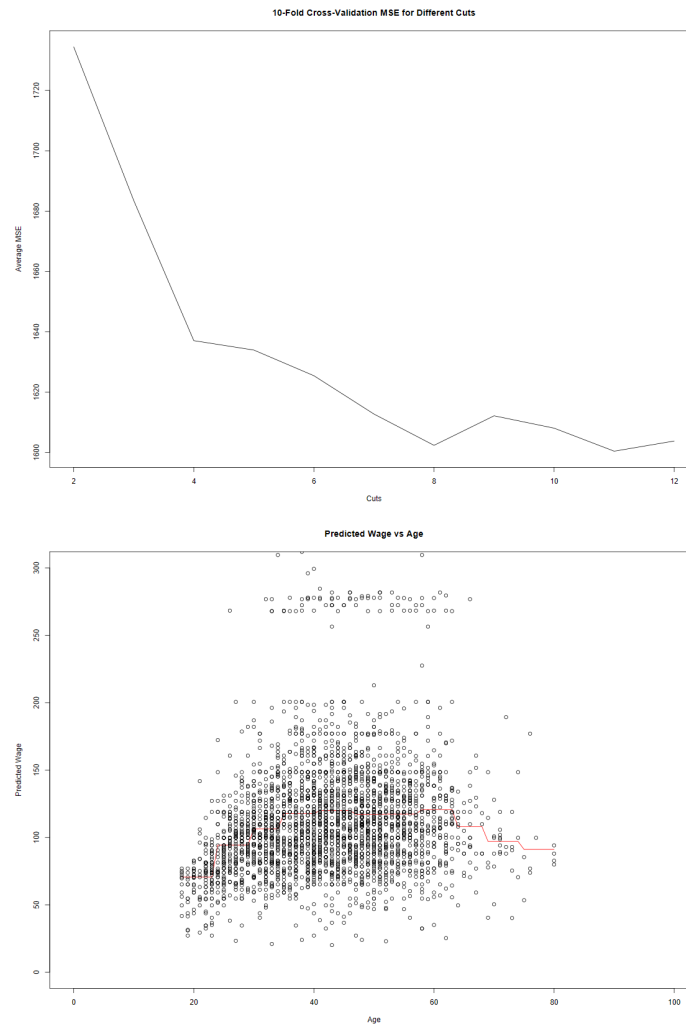


Figure 10

```
library(ISLR2)

data(Wage)
min(Wage$age)
max(Wage$age)

fit_step_function <- function(x, y, k) {
  breaks <- seq(min(x), max(x), length.out = k + 1)

  intervals <- cut(x, breaks = breaks, include.lowest = TRUE)

  step_heights <- tapply(y, intervals, mean)

  predict_step <- function(new_x) {
    new_intervals <- cut(new_x, breaks = breaks, include.lowest = TRUE)
    return(step_heights[as.character(new_intervals)])
  }
}
```

```

return(list(
  breaks = breaks,
  heights = step_heights,
  predict = predict_step
))
}

cv.step <- function(k = 10, Wage, cuts) {
  # get fixed indices for folds and shuffle
  indices <- c(1:nrow(Wage))
  indices <- sample(indices)

  # split into k folds
  folds <- split(indices, rep(1:k, length.out = length(indices)))

  avg_mse <- c()
  for (cut in cuts) {
    mse <- c()
    for (fold in folds) {
      fit <- fit_step_function(Wage$age[-fold], Wage$wage[-fold], cut)
      pred <- fit$predict(Wage[fold, ]$age)
      mse <- c(mse, mean((pred - Wage[fold, ]$wage)^2))
    }
    avg_mse <- c(avg_mse, mean(mse))
  }

  # find the best index
  best_index <- which.min(avg_mse)
  return(list(best_cuts = cuts[best_index], avg_mse = avg_mse, cuts = cuts))
}

cv_results <- cv.step(k = 10, Wage, cuts = 2:12)
cv_results$best_cuts
# best degree is 9

plot(cv_results$avg_mse ~ cv_results$cuts, type = "l", xlab = "Cuts", ylab = "Average MSE", main = "10-Fold C")

# best model
best_model <- fit_step_function(Wage$age, Wage$wage, cv_results$best_cuts)

pred <- best_model$predict(seq(0, 100, 1))

# xlim from 0 to 100
# ylim from 0 to 300
plot(pred ~ seq(0, 100, 1), xlab = "Age", ylab = "Predicted Wage", main = "Predicted Wage vs Age", type = "l",
points(Wage$age, Wage$wage))

models <- lapply(1:12, function(deg) lm(wage ~ poly(age, degree = deg), data = Wage))
# anova all models
do.call(anova, models)

```

The best number of cuts is 11

Applied 8

```
library(ISLR2)
library(splines)

pairs(Auto)

# sort the data by acceleration
Auto <- Auto[order(Auto$acceleration), ]

plot(mpg ~ acceleration, data = Auto)

fit_lm <- lm(mpg ~ acceleration, data = Auto)

# fit a bs spline to the data
fit_bs <- lm(mpg ~ bs(acceleration, df = 4), data = Auto)

# fit a natural spline to the data
fit_ns <- lm(mpg ~ ns(acceleration, df = 4), data = Auto)

# fit a polynomial to the data
fit_poly <- lm(mpg ~ poly(acceleration, degree = 4), data = Auto)

# plot the data
par(mfrow = c(1, 1))
plot(mpg ~ acceleration, data = Auto)

# plot the fitted lines
lines(Auto$acceleration, predict(fit_bs), col = "red")
lines(Auto$acceleration, predict(fit_ns), col = "blue")
lines(Auto$acceleration, predict(fit_poly), col = "green")

# legend
legend("topright", legend = c("bs df = 4", "ns df = 4", "poly df = 4"), col = c("red", "blue", "green"), lty

# normality plot
par(mfrow = c(2, 1))
plot(fit_lm, which = 1)
plot(fit_lm, which = 2)
```

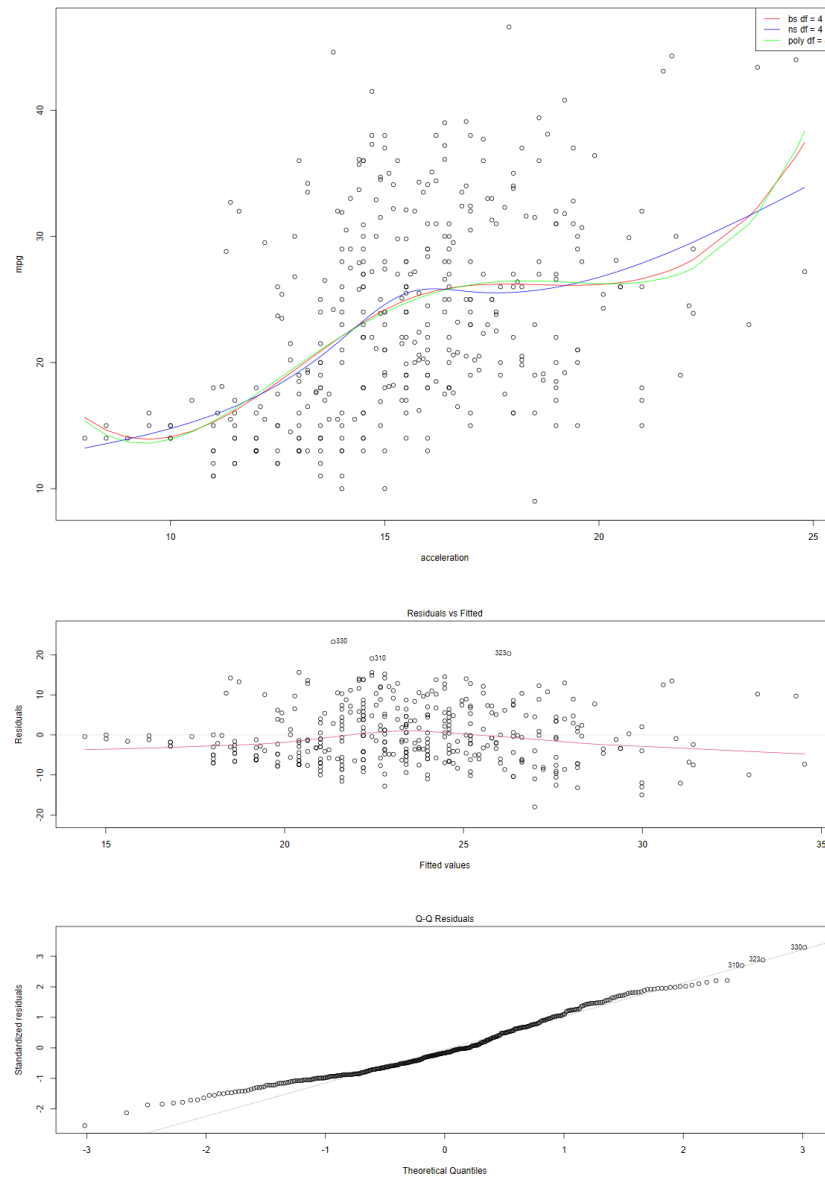


Figure 11

The normality plot shows that the some residuals are not normally distributed.

Applied 9

```

library(ISLR)

str(Boston)
# sort by dis
Boston <- Boston[order(Boston$dis), ]

# fit polynomial regression
fit_poly <- lm(nox ~ poly(dis, degree = 3), data = Boston)
summary(fit_poly)
par(mfrow = c(1, 1))
plot(Boston$dis, Boston$nox)
lines(Boston$dis, predict(fit_poly, Boston), col = "red")
title("Polynomial fit with 3 degrees of freedom")

# fit 1 to 10 degree polynomials

color_map <- rainbow(10)
legend_text <- vector(mode = "character", length = 10)

plot(Boston$dis, Boston$nox)
for (deg in 1:10) {
  fit <- lm(nox ~ poly(dis, degree = deg), data = Boston)
  ssr <- sum(summary(fit)$residuals^2)
  color <- color_map[deg]
  lines(Boston$dis, predict(fit, Boston), col = color)
  legend_text[deg] <- paste("Degree", deg, "SSR:", ssr)
}
legend_text
legend("topright", legend = legend_text, col = color_map, lty = 1)
title("Polynomial fits with degrees 1 to 10")

cv.poly <- function(k = 10, df, degs) {
  # get fixed indices for folds and shuffle
  indices <- c(1:nrow(df))
  indices <- sample(indices)

  # split into k folds
  folds <- split(indices, rep(1:k, length.out = length(indices)))

  avg_mse <- c()
  for (deg in degs) {
    mse <- c()
    for (fold in folds) {
      fit <- lm(nox ~ poly(dis, degree = deg), data = df[-fold, ])
      pred <- predict(fit, df[fold, ])
      mse <- c(mse, mean((pred - df[fold, ]$nox)^2))
    }
    avg_mse <- c(avg_mse, mean(mse))
  }

  # find the best index
  best_index <- which.min(avg_mse)
  return(list(best_deg = degs[best_index], avg_mse = avg_mse, degs = degs))
}

```

```

}

cv_results <- cv.poly(k = 10, df = Boston, degs = 1:13)
cv_results$best_deg

fit.bs <- lm(nox ~ bs(dis, df = 4, knots = c(3)), data = Boston)
plot(Boston$dis, Boston$nox)
lines(Boston$dis, predict(fit.bs, Boston), col = "red")
title("B-spline fit with 4 degrees of freedom and knots at 3")

# Knot is chosen by observing a change of curvature in the plot of nox vs dis

range_of_df <- c(3:6)
par(mfrow = c(2, 2))
for (df in range_of_df) {
  fit <- lm(nox ~ bs(dis, df = df), data = Boston)
  ssr <- sum(summary(fit)$residuals^2)
  plot(Boston$dis, Boston$nox)
  lines(Boston$dis, predict(fit, Boston), col = "red")
  title(paste("B-spline fit with", df, "degrees of freedom and knots at 3", "SSR:", ssr))
}

cv.bs <- function(k = 10, df, degs) {
  # get fixed indices for folds and shuffle
  indices <- c(1:nrow(df))
  indices <- sample(indices)

  # split into k folds
  folds <- split(indices, rep(1:k, length.out = length(indices)))

  avg_mse <- c()
  for (deg in degs) {
    mse <- c()
    for (fold in folds) {
      fit <- lm(nox ~ bs(dis, df = deg), data = df[-fold, ])
      pred <- predict(fit, df[fold, ])
      mse <- c(mse, mean((pred - df[fold, ]$nox)^2))
    }
    avg_mse <- c(avg_mse, mean(mse))
  }

  # find the best index
  best_index <- which.min(avg_mse)
  return(list(best_deg = degs[best_index], avg_mse = avg_mse, degs = degs))
}

cv_results_bs <- cv.bs(k = 10, df = Boston, degs = 4:10)
cv_results_bs$best_deg

```

a, b, c

```
lm(formula = nox ~ poly(dis, degree = 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)	
(Intercept)	0.554695	0.002759	201.021	< 2e-16	***
poly(dis, degree = 3)1	-2.003096	0.062071	-32.271	< 2e-16	***
poly(dis, degree = 3)2	0.856330	0.062071	13.796	< 2e-16	***
poly(dis, degree = 3)3	-0.318049	0.062071	-5.124	4.27e-07	***

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

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

Figure 12

The best degree is 3 from 10 folds cross validation with mse as the metric.

d

The knots is chosen at 3 because it is where the curvature of the plot changes.

e, f

Higher degrees of freedom leads to overfitting. the change of curvature near $x=2$ for $df \geq 4$ does not reflect the true trend.

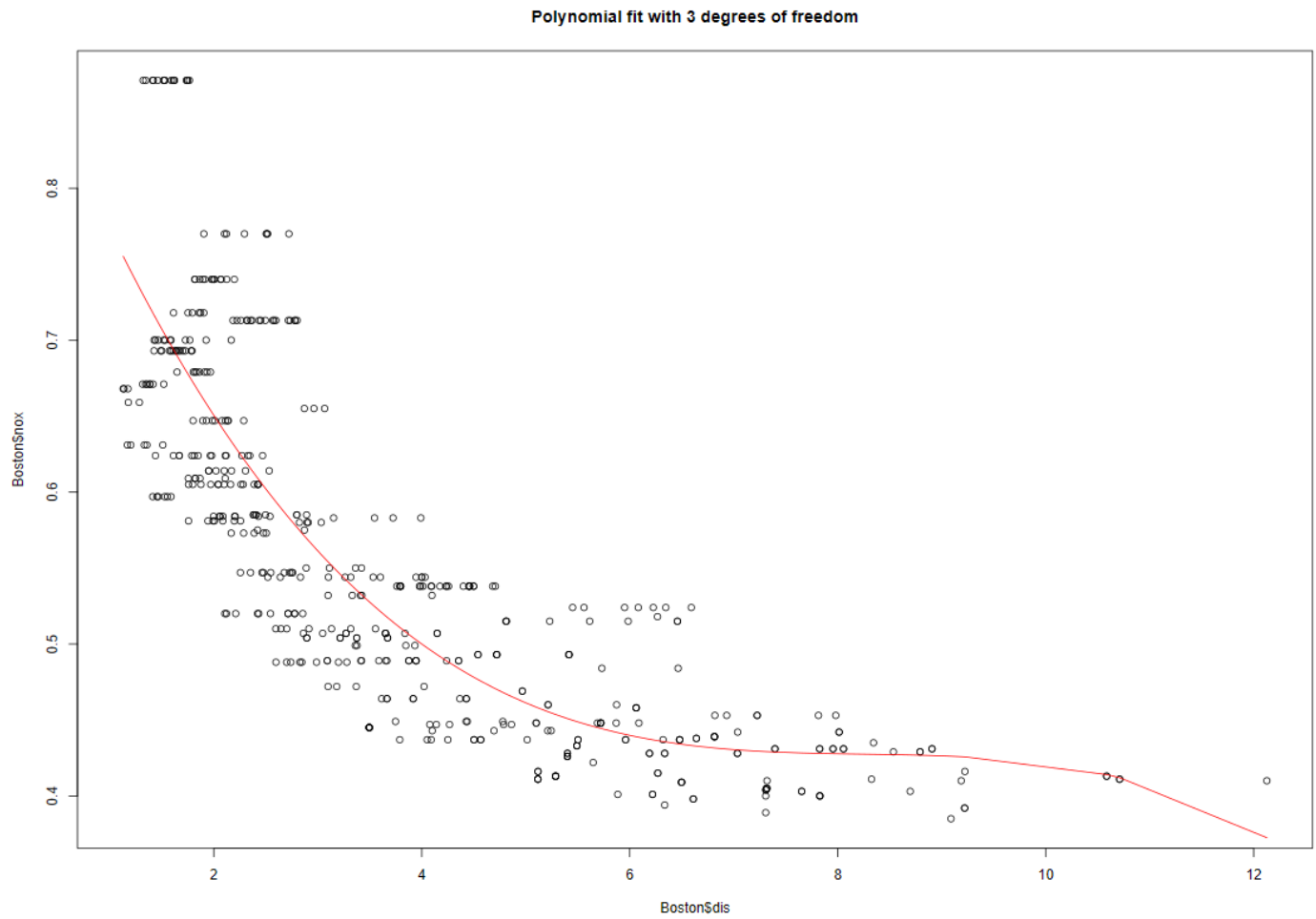


Figure 13

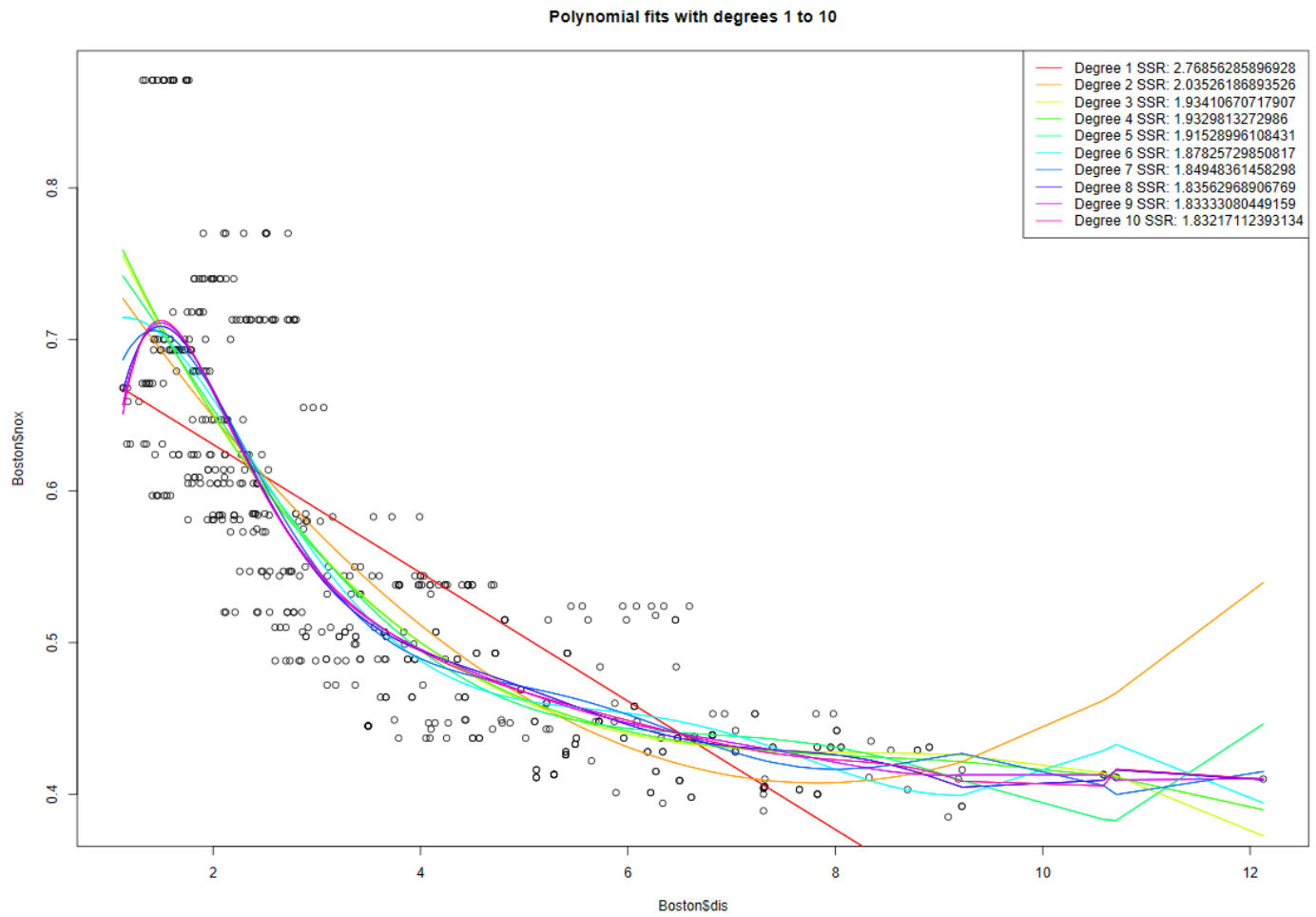


Figure 14

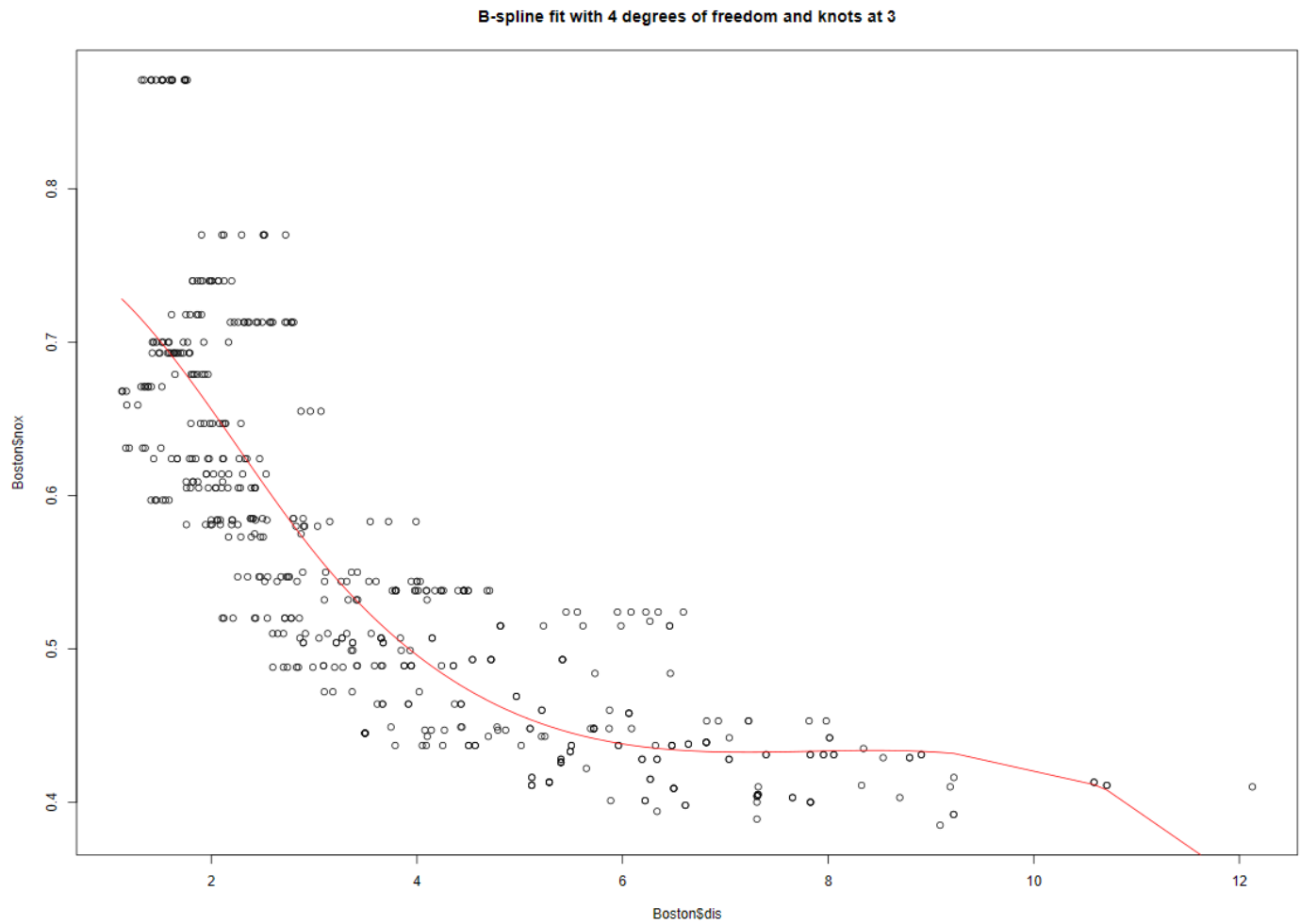


Figure 15

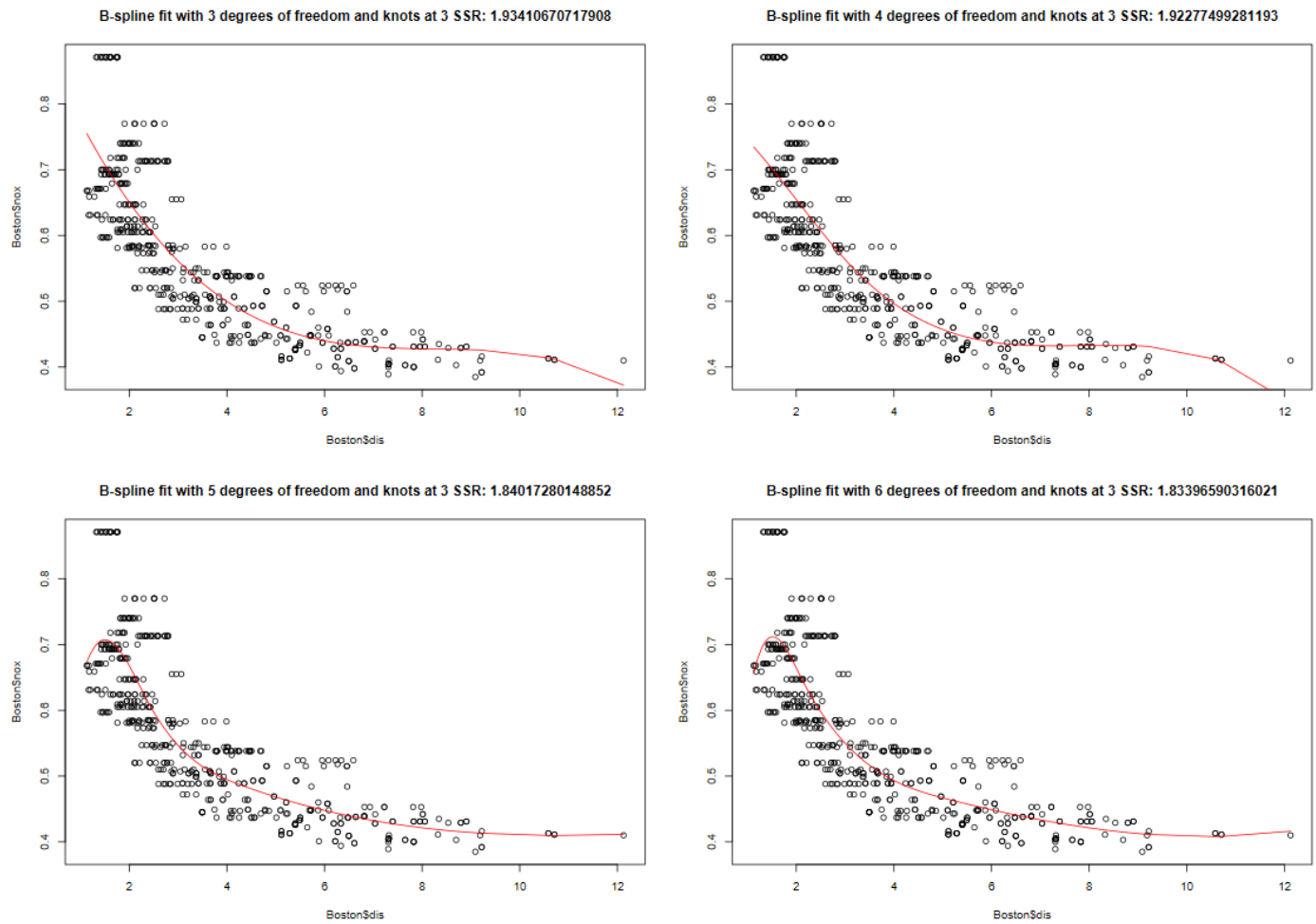


Figure 16