

Regression III

Splines

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Goals for Today

1. Develop the idea of splines as piecewise regression functions.
 - Motivate with piecewise linear model.
2. Consider truncated power basis functions for cubic regression splines.
 - How and where should knots be placed?
3. Discuss other more robust bases - B-splines.
4. Show an example of regression splines at work.

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Definition of Splines

Splines are:

... piecewise regression functions we constrain to join at points called knots (Keele 2007, 70)

- In their simplest form, they are dummy regressors that we use to force the regression line to change direction at some value(s) of X .
- These are similar in spirit to LPR models where we use a subset of data to fit local regressions (but the window doesn't move here).
- These are also allowed to take any particular functional form, but they are a bit more constrained than the LPR model.

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Splines vs. LPR Models

Splines provide a better MSE fit to the data.

- Where $MSE(\hat{\theta}) = \text{Var}(\hat{\theta}) + \text{Bias}(\hat{\theta}, \theta)^2$
- Generally, LPR models will have smaller bias, but much greater variance.
- Splines can be designed to prevent over-fitting (smoothing splines)
- Splines are more easily incorporated in *semi*-parametric models.

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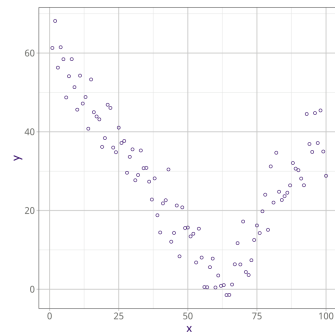
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Regression Splines

We start with the following familiar model:

$$y = f(x) + \varepsilon$$

Here, we would like to estimate this with one model rather than a series of local models.



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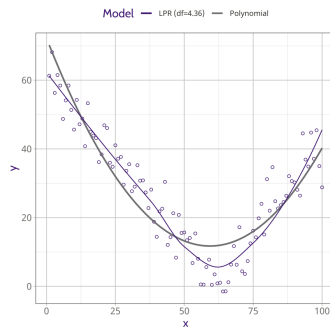
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Failure of Polynomials and LPR

Given what we already learned, we could fit a quadratic polynomial or a LPR:



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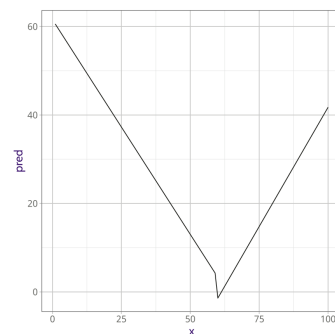
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Dummy Interaction

You might ask, couldn't we just use an interaction between x and a dummy variable coded 1 if $x > 60$ and zero otherwise.

$$y = b_0 + b_1x_1 + b_2d + b_3x \times d + e$$

This seems like a perfectly reasonable thing to do. What can it give you though:



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Basis Functions

A basis function is really just a function that transforms the values of X . So, instead of estimating:

$$y_i = \beta_0 + \beta_1 x_i + \varepsilon_i$$

we estimate:

$$y_i = \beta_0 + \beta_1 b_1(x_i) + \beta_2 b_2(x_i) + \dots + \beta_k b_k(x_i) + \varepsilon_i$$

The basis functions $b_k(\cdot)$ are known ahead of time (not estimated by the model).

- We can think of polynomials as basis functions where $b_j(x_i) = x_i^j$

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Piecewise Polynomials

One way that we can think about regression splines is as piecewise polynomial functions:

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

Just as above though, these polynomials are unconstrained and can generate a discontinuity at the *knot* location c .

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Constraining the Model

To constrain the model, the splines are constructed:

- such that the first and second derivatives of the function continuous.
- Each constraint reduces the number of degrees of freedom we use by one.
- In general, the model uses: Polynomial Degree + # Knots + 1 (for the intercept) degrees of freedom

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Notes

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Truncated Power Basis Functions

The easiest set of Spline functions to consider (for knot location k) are called truncated power functions, defined as:

$$h(x, k) = (x - k)_+^3 = \begin{cases} (x - k)^3 & \text{if } x > k \\ 0 & \text{otherwise} \end{cases}$$

When using these basis functions in, we put the full (i.e., global) parametric function in and a truncated power function of degree n for each knot.

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Linear Truncated Power Functions

To use the truncated power basis for our problem, we need:

- The global linear model
- One truncated power function for the x values greater than the knot location (60).

$$y = b_0 + b_1x + b_2(x - 60)_+^1 + e$$

This sets up essentially 2 equations:

$$x \leq 60 : y = b_0 + b_1x$$

$$x > 60 : y = b_0 + b_1x + b_2(x - 60) = (b_0 - 60b_2) + (b_1 + b_2)x$$

Notice that here we are only estimating 3 parameters, where the interaction would estimate 4 parameters. Thus, this is a constrained version of the interaction.

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Notes

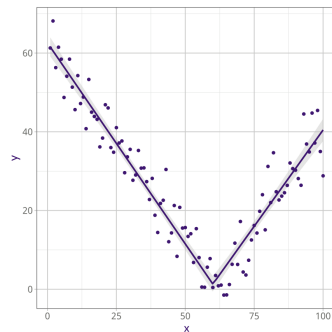
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Fixing the Discontinuity

Including x and $(x - 60)_+$ as regressors, which generates the following predictions:

```
pwl <- function(x, k){ifelse(x >= k, x-k, 0)}  
ggplot(mapping=aes(x=x, y=y)) +  
  geom_smooth(method="lm",  
              formula=y ~ x + pwl(x, 60)) +  
  geom_point() +  
  theme_bw() +  
  mytheme()
```



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Notes

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Polity Example

Thinking back to the Polity example from Lecture 4. We suggested we maybe could fit a piecewise polynomial model:

```
library(foreign)
dat <- read.dta("http://www.quantoid.net/files/reg3/linear_ex.dta")
dat$polity_dem_fac <- as.factor(dat$polity_dem)
unrestricted.mod <- lm(rep1 ~ polity_dem_fac + iwar +
  cwar + logpop + gdppc, data=dat)
pwl.mod <- lm(rep1 ~ polity_dem + pwl(polity_dem, 9) +
  iwar + cwar + logpop + gdppc, data=dat)
anova(pwl.mod, unrestricted.mod, test="F")

## Analysis of Variance Table
##
## Model 1: rep1 ~ polity_dem + pwl(polity_dem, 9) + iwar + cwar + logpop +
##   gdppc
## Model 2: rep1 ~ polity_dem_fac + iwar + cwar + logpop + gdppc
##   Res.Df    RSS Df Sum of Sq    F Pr(>F)
## 1      2676 2172.9
## 2      2668 2163.3   8      9.651 1.4878 0.1562
```



Notes

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Unknown Knot Location

If you don't know the knot location, you could try a bunch of different options.

```
cvfun <- function(split, ...){
  mods <- lapply(1:9, function(i){
    lm(rep1 ~ polity_dem + pwl(polity_dem, i) +
      iwar + cwar + logpop + gdppc,
      data= analysis(split)))
  yhat <- sapply(mods, function(x)predict(x, newdata=assessment(split)))
  y <- assessment(split)$rep1
  e <- apply(yhat, 2, function(z)(y-z)^2)
  sume2 <- colSums(e)
  n <- length(y)
  tibble(knot = 1:9, e2 = sume2, n = rep(n, 9))
}
out <- dat %>%
  vfold_cv(v=10, repeats=3) %>%
  mutate(err = map(splits, cvfun)) %>%
  unnest(err) %>%
  group_by(id, knot) %>%
  summarise(mse = sum(e2)/sum(n)) %>%
  ungroup %>%
  group_by(knot) %>%
  summarise(mse = mean(mse))
```

```
aics <- sapply(1:9, function(i)
  AIC(lm(rep1 ~ polity_dem + pwl(polity_dem, i) +
    iwar + cwar + logpop + gdppc,
    data= dat)))
out <- out %>%
  mutate(aic = aics)
out

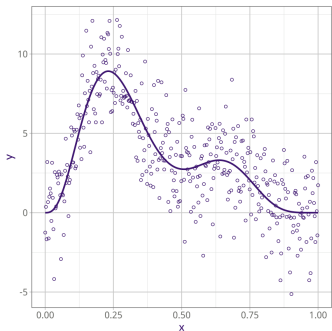
## # A tibble: 9 x 3
##   knot mse aic
##   <int> <dbl> <dbl>
## 1     1  0.934 7431.
## 2     2  0.929 7417.
## 3     3  0.919 7387.
## 4     4  0.907 7354.
## 5     5  0.900 7331.
## 6     6  0.887 7293.
## 7     7  0.868 7235.
## 8     8  0.840 7145.
## 9     9  0.815 7064.
```

Notes

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Example: Cubic Spline

Consider the following relationship:



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Cubic Spline

$$y = b_0 + b_1x + b_2x^2 + b_3x^3 + \sum_{m=1}^{\# \text{ knots}} b_{k+3}(x - k_m)_+^3$$

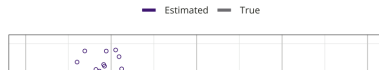
Let's consider our example with 3 knots $k = \{.2, .4, .6, .8\}$

```
## Coefficients:
##      Estimate Std. Error t value Pr(>|t|)
## (Intercept)      8.602e-02  6.836e-01   0.126   0.8999
## x             -3.885e+00  1.894e+01  -0.205   0.8376
## I(x^2)          5.772e+02  1.386e+02   4.164 3.84e-05 ***
## I(x^3)         -1.783e+03  2.877e+02  -5.921 6.99e-09 ***
## I((x - k[1])^3 * (x >= k[1])) 2.771e+03  3.789e+02   7.314 1.48e-12 ***
## I((x - k[2])^3 * (x >= k[2])) -1.474e+03  1.821e+02  -8.094 7.36e-15 ***
## I((x - k[3])^3 * (x >= k[3]))  3.866e+02  1.821e+02   2.123  0.0344 *
## I((x - k[4])^3 * (x >= k[4]))  7.888e+02  3.789e+02   1.869  0.0624 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard deviation: 1.951 on 392 degrees of freedom
## Multiple R-squared:  0.6665
## F-statistic: 111.9 on 7 and 392 DF,  p-value: < 2.2e-16
##      AIC      BIC
## 1679.71 1715.63
```

Notes

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Predictions



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Problems with Truncated Power Basis Functions

- Highly collinear and can lead to instability and singularities (i.e., computationally bad stuff) at worst.
- Not as "local" as some other options, the support of the piecewise functions can be over the whole range of the data or nearly the whole range of the data.
- Can produce erratic tail behavior.

Other basis functions, like the B-spline basis functions solve all of these problems:

- Reduces collinearity (though doesn't eliminate it)
- Support of the function is more narrowly bounded.
- Uses knots at the boundaries of x and assumes linearity beyond the knots.

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Notes

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Example: B-spline

```
library(splines)
k <- c(.2,.4,.6,.8)
csmod2 <- lm(y ~ bs(x, knots=k))
S(csmod2, brief=TRUE)

## Coefficients:
##           Estimate Std. Error t value Pr(>|t|)
## (Intercept)    0.08602     0.68358   0.126  0.8999
## bs(x, knots = k)1 -0.25898     1.26296  -0.205  0.8376
## bs(x, knots = k)2 14.61385     0.80729 18.102 < 2e-16 ***
## bs(x, knots = k)3  1.33876     0.90742   1.384  0.1672
## bs(x, knots = k)4  0.83755     4.46310  0.186  0.8585
## bs(x, knots = k)5  2.30614     1.01655   2.269  0.0238 *
## bs(x, knots = k)6 -1.83334     0.99321  -1.846  0.0657 .
## bs(x, knots = k)7  0.80657     0.97507   0.827  0.4086
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard deviation: 1.951 on 392 degrees of freedom
## Multiple R-squared: 0.6665
## F-statistic: 111.9 on 7 and 392 DF,  p-value: < 2.2e-16
##           AIC          BIC
## 1679.71 1715.63
```

Notice that the fit here is precisely the same as with the truncated power basis functions

```
p1 <- predict(csmod, se=TRUE)
p2 <- predict(csmod2, se=TRUE)
cor(p1$fit, p2$fit)

## [1] 1

cor(p1$se.fit, p2$se.fit)

## [1] 1
```

Notes

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Interpreting Spline Coefficients

So, how do you interpret the spline coefficients?

- You don't.
- Remember that these are all functions of x , so we cannot change the values of one component of the basis function while holding the others constant, the others would have to change, too.

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Choices in Spline Models

- **Degree:** the analyst has to choose the degree of the polynomial fit to the subsets of the data.
- **Number of knots:** the analyst has to choose the number of knots
- **Location of knots:** Often, knots are spaced evenly over the support of the data (i.e., the range of x), but that needn't be the case.
 - Knot placement can be guided by theory if possible.
 - Otherwise, for the functions we generally need to estimate, a few knots should probably work just fine.

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Notes

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How important is knot placement?

I did a simulation where I did the following:

- Using the function created above, I first estimated a B -spline with 1-10 knots.
 - Then, I calculated the R^2 with respect to the true point location.
- For each number of knots, randomly draw knots from a uniform distribution.
 - Estimate the model and calculate R^2 with respect to the truth.

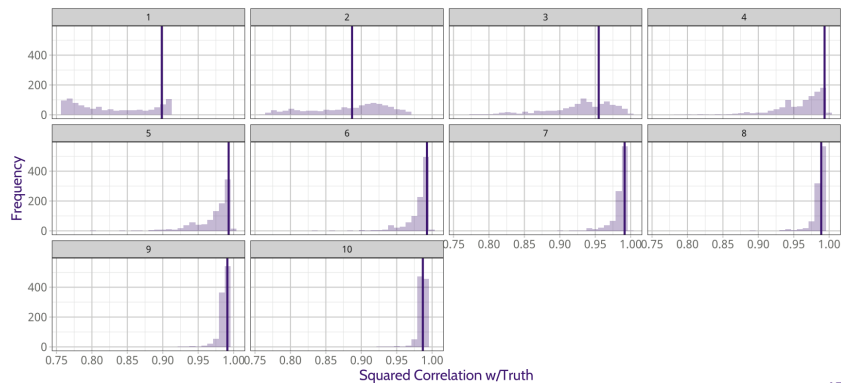
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Notes

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Results



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How Important is Knot Placement? II

- So long as the polynomial degree is reasonably high (3 should be high enough for what we do, but 4 might be useful if you have a very complicated function), knot placement is not particularly important.
- Use theory, if it exists, to place knots.
- If theory doesn't exist, knots placed evenly across the range of x will, in general, minimize error.
- If you think about the knots as random variables (because we don't know their values) and further that they are distributed uniformly (i.e., neither middle or extreme values are more likely), then technically evenly spaced knots minimize distance to the true, but unknown knots.

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How Important is Polynomial Degree?

- Pretty important, particularly if we don't know or have a really good sense of where the knots should be.
- B-splines are more forgiving of knot placement errors the higher the polynomial degree.
- Generally no good reason to use something more restrictive than a cubic spline.
- We are generally not trying to model particularly complicated functions.
- More knots are more likely to be used than a higher polynomial degree to make the function more flexible.

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How Important is the Number of Knots

- Flexibility increases with number of knots and polynomial degree.
- Increasing number of knots can make the function more flexible.
- We can use AIC, BIC or Cross-Validation to choose number of knots.

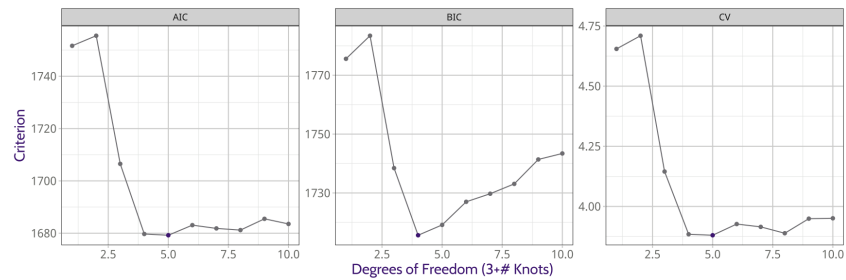
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Notes

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Choosing Number of Knots



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Notes

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Worked Example

```
library(car)
library(rio)
dat <- import(
  "http://www.quantoid.net/files/reg3/jacob.dta")
rawlm <- lm(chal_vote ~ perotvote + chal_spend +
  exp_chal, data=dat)
```

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Notes

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Raw Model

```
S(rawlm, brief=TRUE)

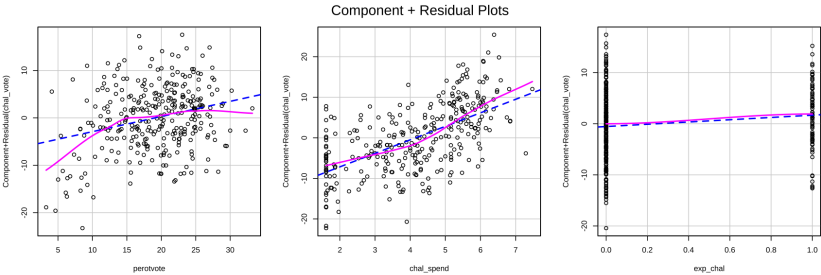
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 15.95365    1.59783   9.990 < 2e-16 ***
## perotvote   0.31943    0.06655   4.800 2.48e-06 ***
## chal_spend  3.33294    0.27869  11.959 < 2e-16 ***
## exp_chal    2.22853    0.98576   2.253  0.025 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard deviation: 6.74 on 308 degrees of freedom
## Multiple R-squared: 0.4552
## F-statistic: 85.8 on 3 and 308 DF, p-value: < 2.2e-16
##      AIC      BIC
## 2082.02 2100.74
```

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C+R Plots

```
crPlots(rawlm, layout=c(1,3))
```



Notes

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Transformation Model

```
boxTidwell(chal_vote ~ perotvote,
            ~ chal_spend + exp_chal, data=dat)

## MLE of lambda Score Statistic (z) Pr(>|z|)
## ---
## -1.0634 -4.1129 3.908e-05 ***
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## iterations = 9

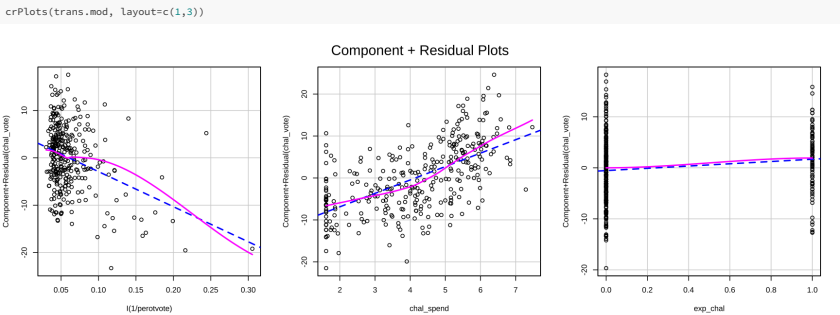
trans.mod <- lm(chal_vote ~ I(1/perotvote) +
                chal_spend + exp_chal, data=dat)
S(trans.mod, brief=TRUE)

## Coefficients:
## Estimate Std. Error t value Pr(>|t|)
## (Intercept) 27.0997 1.4276 18.983 < 2e-16 ***
## I(1/perotvote) -74.0400 11.0670 -6.346 7.9e-10 ***
## chal_spend 3.1989 0.2737 11.687 < 2e-16 ***
## exp_chal 2.2218 0.9610 2.312 0.0214 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard deviation: 6.571 on 308 degrees of freedom
## Multiple R-squared: 0.4822
## F-statistic: 95.6 on 3 and 308 DF, p-value: < 2.2e-16
## AIC BIC
```

Notes

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C+R Plots

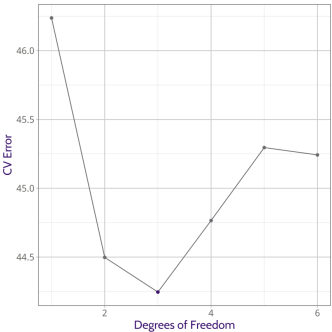


Notes

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Degrees of Freedom

```
library(DAHAnc)
nkp <- NKnotes(chal_vote ~ chal_spend + exp_chal,
  "perotvote", max.knots=3, data=dat, includePol
  criterion="CV", plot=FALSE, cviter=10)
nkp$df <- 1:6
nkp$min <- factor(ifelse(nkp$stat == min(nkp$stat),
  levels=c(0,1),
  labels=c("Other", "Minimum")))
```



Notes

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Polynomial

```
poly.mod <- lm(chal_vote ~ poly(perotvote, 3) +
  chal_spend + exp_chal, data=dat)
S(poly.mod, brief=TRUE)
```

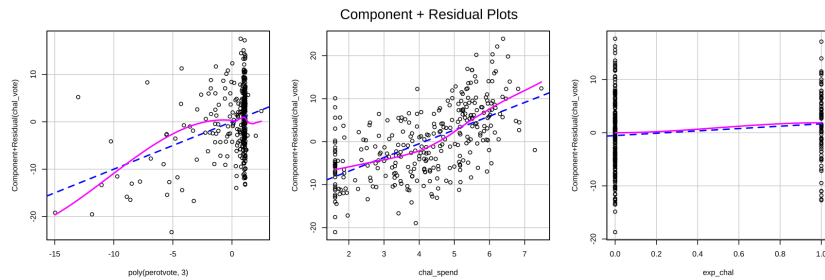
```
## Coefficients:
##           Estimate Std. Error t value Pr(>|t|)
## (Intercept)    22.6040      1.1222   20.143 < 2e-16 ***
## poly(perotvote, 3)1    33.2168      6.6421    5.001 9.63e-07 ***
## poly(perotvote, 3)2   -25.5074      6.5992   -3.865 0.000136 ***
## poly(perotvote, 3)3    11.8753      6.6112    1.796 0.073444 .
##   chal_spend      3.2801      0.2738   11.688 < 2e-16 ***
##   exp_chal       2.2016      0.9613    2.290 0.022683 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard deviation: 6.571 on 306 degrees of freedom
## Multiple R-squared:  0.4856
## F-statistic: 57.76 on 5 and 306 DF,  p-value: < 2.2e-16
##           AIC           BIC
## 2068.16 2094.36
```

Notes

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More CR Plots

```
crPlots(poly.mod, layout=c(1,3))
```



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Notes

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Which is better?

```
library(clarkeTest)
clarke_test(trans.mod, poly.mod)
```

```
##
## Clarke test for non-nested models
##
## Model 1 log-likelihood: -1028
## Model 2 log-likelihood: -1027
## Observations: 312
## Test statistic: 207 (66%)
##
## Model 1 is preferred (p = .8e-09)
```

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Notes

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Challenger Spending

```
boxTidwell(chal_vote ~ chal_spend,
~ I(1/perotvote) + exp_chal,
data=dat)

## MLE of lambda Score Statistic (z) Pr(>|z|)
## 2.2987 3.7127 0.0002051 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## iterations = 5

boxTidwell(chal_vote ~ I(chal_spend^2),
~ I(1/perotvote) + exp_chal,
data=dat)

## MLE of lambda Score Statistic (z) Pr(>|z|)
## 1.1495 0.8603 0.3896
##
## iterations = 4

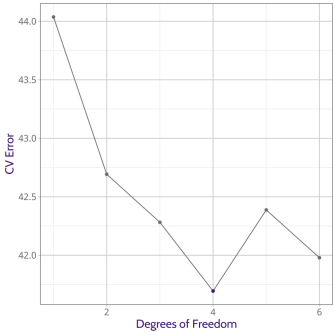
trans.mod2 <- lm(chal_vote ~I(chal_spend^2) +
I(1/perotvote) + exp_chal, data=dat)
```

Notes

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Degrees of Freedom

```
library(DAHAisc)
nkp <- NKnotes(chal_vote ~ I(1/perotvote) + exp_c
"chal_spend", max.knots=3, data=dat, includePo
criterion="CV", plot=FALSE, cviter=10)
nkp$ddf <- 1:6
nkp$min <- factor(ifelse(nkp$stat == min(nkp$sta
levels=c(0,1),
labels=c("Other", "Minimum")))
```



Notes

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Spline Model

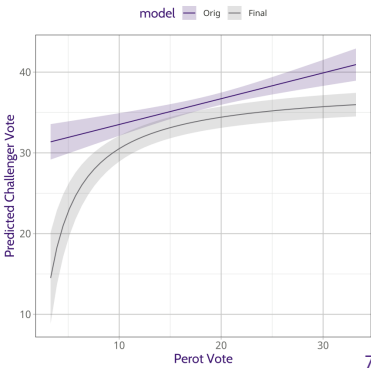
```
spline.mod <- dat %>%
  mutate(inv_perotvote = 1/perotvote) %>%
  lm(chal_vote ~ inv_perotvote +
    bs(chal_spend, df=4) +
    exp_chal, data=.)
anova(trans.mod2, spline.mod, test="F")

## Analysis of Variance Table
##
## Model 1: chal_vote ~ I(chal_spend^2) + I(1/perotvote) + exp_chal
## Model 2: chal_vote ~ inv_perotvote + bs(chal_spend, df = 4) + exp_chal
## Res.Df  RSS Df Sum of Sq    F    Pr(>F)
## 1    308 12816
## 2    305 12208  3    608.53 5.0679 0.001939 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Notes

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Effects



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Redux

Splines can be a good way to ...

- Fit models that cannot be easily fit by other simpler parametric forms still within the OLS/GLM framework.
- Test a wider array of possible alternative functional forms.

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Notes

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