# Model Diagnostics

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Regression Analysis

- ▶ Residuals *ei* vs.
  - ► *i* (independence)

  - *x*<sub>1</sub>, ..., *x*<sub>k</sub> (linearity)
    Ŷ<sub>i</sub> (linearity and homogeneity of variance)

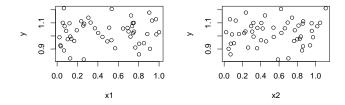
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  - $ightharpoonup \hat{Y}_i$  (linearity and homogeneity of variance)
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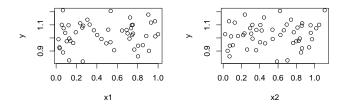
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- Now we'll discuss
  - added variable plots,
  - leverages,
  - DFFITS,
  - Cook's distance.

# The problem with marginal plots

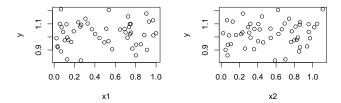


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(Intercept) 0.05083 0.07726 0.658 0.514
x1 0.95536 0.07651 12.487 <2e-16 ***
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Outliers, leverages, DFFITs, Cook's distance

Review of Diagnostics

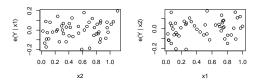
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where  $e(Y \mid x_i)$  are residuals when we regress Y on  $x_i$  only.



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- ▶ An added variable plot tries to fix this problem.
- ▶ It answers the question: Does  $x_j$  explain any *residual* variability once the rest of the predictors are in the model?

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- ▶ If you fit a simple linear regression

$$e_i(Y \mid \mathbf{x}_{-j}) = \beta_j \cdot e_i(x_j \mid \mathbf{x}_{-j}) + \epsilon_i$$

then the LSE  $\hat{\beta}_j$  is the same as one would get from fitting the full model  $Y_i = \beta_0 + \beta_1 x_{i1} + \cdots + \beta_k x_{ik} + \epsilon_i$ .

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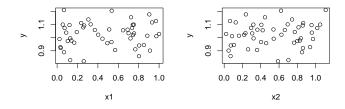
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▶ Gives an idea of the functional form of  $x_j$ : a transformation in  $x_j$  should mimic the pattern seen in the plot.



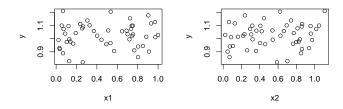
#### 10.1 Added variables plots illustration

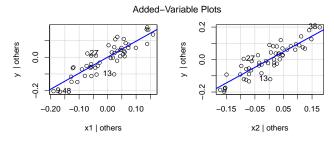
Let's go back to our synthetic example:



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Let's go back to our synthetic example:



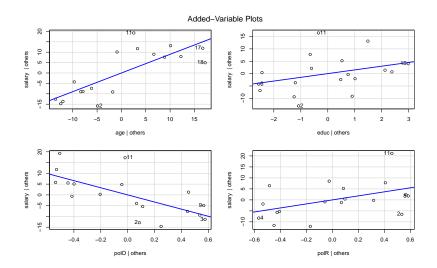


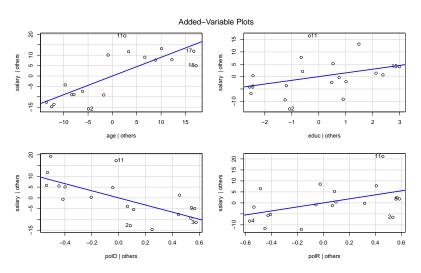
# Salary data, first order terms only

#### Coefficients:

```
Estimate Std. Error t value Pr(>|t|)
(Intercept) 17.0313 7.3459 2.318 0.03735 *
age 0.8983 0.1968 4.565 0.00053 ***
educ 1.5039 1.1841 1.270 0.22632
pol0 -16.5404 4.8807 -3.389 0.00484 **
polR 9.1587 4.8482 1.889 0.08139 .
```

Residual standard error: 8.209 on 13 degrees of freedom Multiple R-squared: 0.8374, Adjusted R-squared: 0.7873





Age effect is nonlinear; let's add a quadratic term.

# Salary data, quadratic effect in age

```
> m1 = lm(salary ~ . + I(age^2), data=salary_data)
> summary(m1)
```

#### Coefficients:

```
Estimate Std. Error t value Pr(>|t|)

(Intercept) -39.224169 16.810142 -2.333 0.037839 *

age 3.463723 0.740666 4.676 0.000535 ***

educ 2.166475 0.883369 2.453 0.030453 *

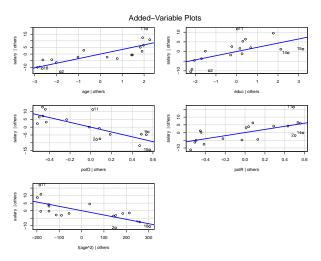
pol0 -15.455108 3.571147 -4.328 0.000983 ***

polR 10.118144 3.544586 2.855 0.014500 *

I(age^2) -0.028831 0.008166 -3.530 0.004143 ***
```

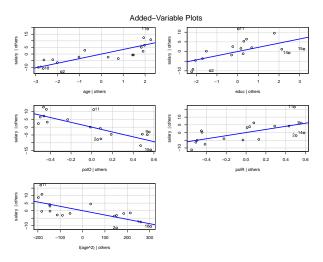
Residual standard error: 5.984 on 12 degrees of freedom Multiple R-squared: 0.9202, Adjusted R-squared: 0.887

# Added variables plots w/ quadratic age



Education is now significant!

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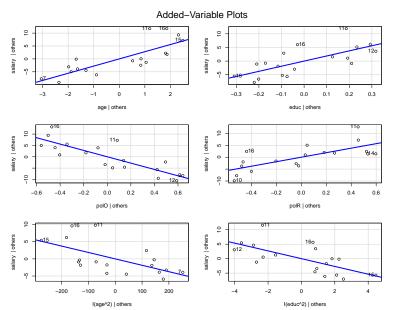
Education is now significant! The incorrect functional form for age was *masking* the importance of education.

## Salary data, quadratic effect in education

```
> summary(m2)
Call:
lm(formula = salary ~ . + I(age^2) + I(educ^2), data = salary_data)
            Estimate Std. Error t value Pr(>|t|)
(Intercept) -75.977348 18.262402 -4.160 0.001589 **
age
    2.787032 0.626151 4.451 0.000977 ***
educ 18.751324 5.739109 3.267 0.007501 **
pol0 -13.976910 2.848879 -4.906 0.000467 ***
polR 9.495127 2.790631 3.403 0.005903 **
I(age^2) -0.018677 0.007298 -2.559 0.026558 *
I(educ^2) -1.342341 0.461108 -2.911 0.014161 *
```

Residual standard error: 4.697 on 11 degrees of freedom Multiple R-squared: 0.9549, Adjusted R-squared: 0.9304

# Added variable plots w/ quad. age and educ



Outliers, leverages, DFFITs, Cook's distance

Review of Diagnostics

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  - if not, outlying values may skew inference a lot and yield models with poor predictive properties.

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- ▶ **Cook's distance** is a consolidated measure of influence the point  $(\mathbf{x}_i, Y_i)$  has on the regression surface at all n points  $\mathbf{x}_1, \ldots, \mathbf{x}_n$ .



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$$\begin{aligned} \operatorname{Cov}[\hat{\mathbf{Y}}] &= \operatorname{Cov}[\mathbf{X}\hat{\boldsymbol{\beta}}] = \operatorname{Cov}[\mathbf{X}(\mathbf{X}^{\top}\mathbf{X})^{-1}\mathbf{X}^{\top}\mathbf{Y}] \\ &= \operatorname{Cov}[\mathbf{H}\mathbf{Y}] = \mathbf{H}\operatorname{Cov}[\mathbf{Y}]\mathbf{H}^{\top} = \mathbf{H}\sigma^{2}\mathbf{I}\mathbf{H}^{\top} \end{aligned}$$

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 $\Rightarrow \operatorname{Var}[\hat{Y}_i] = \sigma^2 h_{ii}$ , and its unbiased estimator is MSE  $\cdot h_{ii}$ .

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- A refinement of the standardized residual that has a recognizable distribution is the studentized deleted residual

$$t_i = \frac{Y_i - \hat{Y}_i}{\sqrt{\mathsf{MSE}_{(i)}(1 - h_{ii})}}$$

where  $MSE_{i(i)}$  is obtained from the model when *i*-th example was removed from the data.



In fact, no need to fit n additional regressions, because there is relationship b/w MSE and MSE<sub>(i)</sub>:

$$(n-p)\mathsf{MSE} = (n-p-1)\mathsf{MSE}_{(i)} + \frac{e_i^2}{1-h_{ii}}$$
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▶ Typically, in practice, one simply flags observations with  $|t_i| > t_{1-\alpha/2;n-p-1}$  as *possibly* outlying in consideration with other diagnostics.



► The leverages  $h_{ii}$  get larger the further the points  $\mathbf{x}_i$  are from the mean  $\bar{\mathbf{x}} = \frac{1}{n} \sum_{i=1}^{n} \mathbf{x}_i$ , adjusted for "how many" other predictors are in the vicinity of  $\mathbf{x}_i$ .

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- ▶ Use the fact that  $\mathbf{H} = \mathbf{X}(\mathbf{X}^{\top}\mathbf{X})^{-1}\mathbf{X}^{\top} = \mathbf{H}\mathbf{H}$  to show  $\sum_{i=1}^{n} h_{ii} = p$  and  $0 \le h_{ii} \le 1$ .

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- ▶ The rule of thumb is that any leverage  $h_{ii}$  that is larger than twice the mean leverage p/n, i.e.  $h_{ii} > 2p/n$ , is flagged as having "high" leverage.

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- ▶ In R, you can get leverages using hatvalues command.

▶ The  $i^{th}$  DFFIT, denoted DFFIT<sub>i</sub>, is given by

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#### 10.4 Cook's distance

► The *i*<sup>th</sup> Cook's distance, denoted *D<sub>i</sub>*, is an aggregate measure of the influence of the *i*<sup>th</sup> observation on all *n* fitted values:

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► Look for values of Cook's distance significantly larger than other values; these are cases that have disproportionate influence on the fitted regression surface as a whole.

Added variable plots

Outliers, leverages, DFFITs, Cook's distance

Review of Diagnostics

Example

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An alternative approach that allows keeping correlated predictors is ridge regression (Chapter 11).

▶ Deleted residuals  $t_i \sim t_{n-p-1}$ , so you can formally define an outlier as being larger than  $t_{1-\alpha/(2n),n-p-1}$ .

▶ Residual plots. Plots of  $e_i$  or  $t_i$  vs.  $\hat{Y}_i$  and versus each  $x_1, \ldots, x_k$  help assess (a) correct functional form, (b) constant variance, and (c) outlying observations. They may also suggest a transformation for a predictor or two.

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Added variable plots help figure out functional form of predictors, and whether significance is being driven by one or two points only.

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- ► A **normal Q-Q plot** of the residuals will indicate departures from normality.
- ▶ A list of the studentized deleted residuals, leverages, and Cook's distances helps to determine outlying values that may be transcription errors or data anomalies and also indicates those observations that affect the fitted regression surface as a whole.

#### Standard diagnostic plots

- ▶ t<sub>i</sub> vs. h<sub>i</sub>. Which observations are outlying in x-direction, outlying in Y-direction, or both?
- ▶ D<sub>i</sub> vs. i. Which observations grossly affect fit of regression surface?
- $ightharpoonup e_i$  vs.  $\hat{Y}_i$  and  $t_i$  vs.  $\hat{Y}_i$ . Constant variance & linearity.
- ▶  $Y_i$  vs.  $\hat{Y}_i$ ; how well model predicts its own data. Better models have points close to line y = x.
- ▶ Normal probability plot of the  $e_1, ..., e_n$ .
- ▶ Histogram of  $e_1, ..., e_n$ .
- ▶ Plots of  $e_i$  vs. each predictor  $x_1, ..., x_k$ .
- ▶ One more plot that prof. Hanson never looks at.

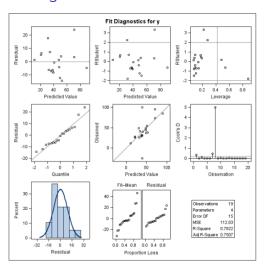
Added variable plots

Outliers, leverages, DFFITs, Cook's distance

Review of Diagnostics

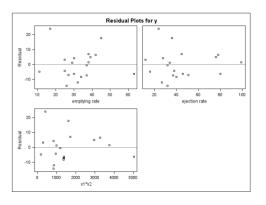
Example

#### An example of diagnostics



Model is  $Y_i = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \beta_{12} x_{i1} x_{i2} + \epsilon_i$ . One highly influential point & one poorly fit.

## Residual plots



These look pretty good, except for the one large residual.

#### Arterial pressure data

```
proc glm;
model y=x1 x2 x1*x2;
output out=out cookd=c rstudent=r; run;
proc print; var x1 x2 y c r; run;
Obs
       x1
                                        r
      45
             36
                   49
                         0.36904
                                     2.20950
                        0.00383
                                     0.39889
            16
                        0.12052
                                    -0.62921
      11
      30
                        0.00885
                                    -0.60493
      39
            76
                        0.01498
                                 0.51721
                        0.02392
                                     0.66178
      17
             24
                        0.45892
                                 3.31414
 8
      63
            80
                   26
                        4.99081
                                    -1.77941
      25
            12
                   74
                        0.00724
                                     0.33794
10
       32
             27
                   37
                         0.04100
                                    -1.22324
11
      37
             37
                         0.01660
                                    -0.71526
                   31
12
                        0.00032
                                   0.12816
13
      26
             32
                   38
                        0.04023
                                    -1.45743
14
                        0.01271
                                  0.69211
15
                        0.00817
                                     0.18206
      25
                        0.00422
                                    -0.40213
16
17
                        0.02196
                                 -0.70921
18
      37
                         0.00014
                                  -0.05730
                   40
19
       34
                   31
                         0.01371
                                    -0.80210
```

Obs. 7 has largest arterial pressure. Obs. 8 has relatively small arterial pressure.

# Dropping obs. 8 and obs. 7

proc glm data=out; model y=x1 x2 x1*x2; run;					
		Standard			
Parameter	Estimate	Error	t Value	Pr >  t	
Intercept	134.3998664	15.98159869	8.41	<.0001	
x1	-2.1330220	0.52215739	-4.09	0.0010	
x2	-1.6993299	0.36366865	-4.67	0.0003	
x1*x2	0.0333471	0.00928281	3.59	0.0027	
proc glm data=out(where=(c<4)); model y=x1 x2 x1*x2; run;					
		Standard			
Parameter	Estimate	Error	t Value	Pr >  t	
Intercept	157.5094488	19.79515582	7.96	<.0001	
x1	-2.7122125	0.58667658	-4.62	0.0004	
x2	-2.7743376	0.69321545	-4.00	0.0013	
x1*x2	0.0618590	0.01822201	3.39	0.0044	
<pre>proc glm data=out(where=(abs(r)&lt;3)); model y=x1 x2 x1*x2; run;</pre>					
		Standard			
Parameter	Estimate	Error	t Value	Pr >  t	
Intercept	116.3928224	13.52293668	8.61	<.0001	
x1	-1.6161083	0.43361763	-3.73	0.0023	
x2	-1.4903775	0.28875668	-5.16	0.0001	
x1*x2	0.0272510	0.00742428	3.67	0.0025	

How do 7 and 8 affect the significance and/or magnitude of the effects?