

The Language of Models

Author: Nicholas G Reich

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Today's topics

- The language of models
- Model formulas and coefficients

Example: predicting respiratory disease severity (“lung” dataset)

Reading: Kaplan, Chapters 6 and 7.



Figure acknowledgements to [Hadley Wickham](#).

Lung Data Example

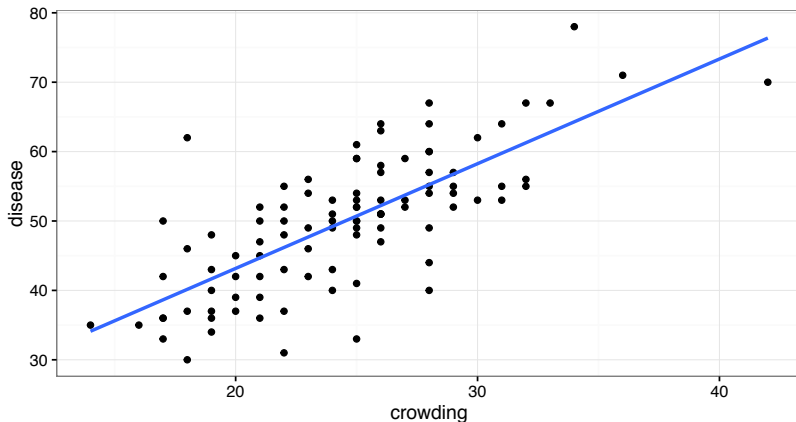
99 observations on patients who have sought treatment for the relief of respiratory disease symptoms.

The variables are:

- `disease` measure of disease severity (larger values indicates more serious condition).
- `education` highest grade completed
- `crowding` measure of crowding of living quarters (larger values indicate more crowding)
- `airqual` measure of air quality at place of residence (larger number indicates poorer quality)
- `nutrition` nutritional status (larger number indicates better nutrition)
- `smoking` smoking status (1 if smoker, 0 if non-smoker)

Lung Data Example: terms defined

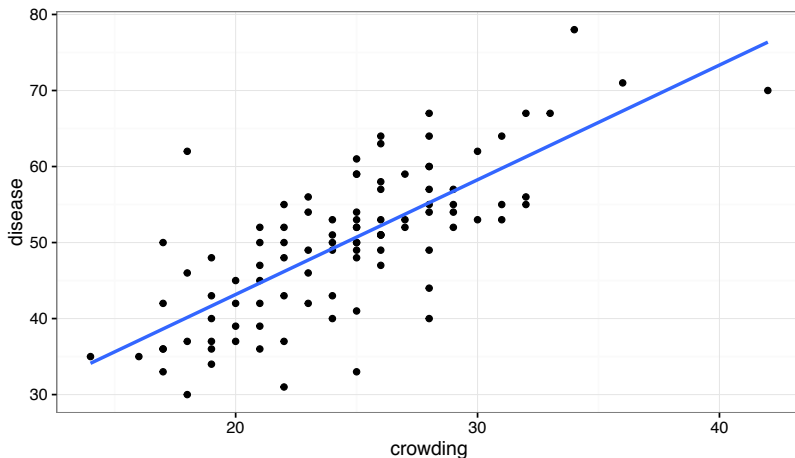
```
dat <- read.table("lungc.txt", header=TRUE)
ggplot(dat, aes(crowding, disease)) + geom_point() +
  geom_smooth(method="lm", se=FALSE)
```



Things to point out: response variable? explanatory variable?
model value? residual?

Lung Data Example: what is the model?

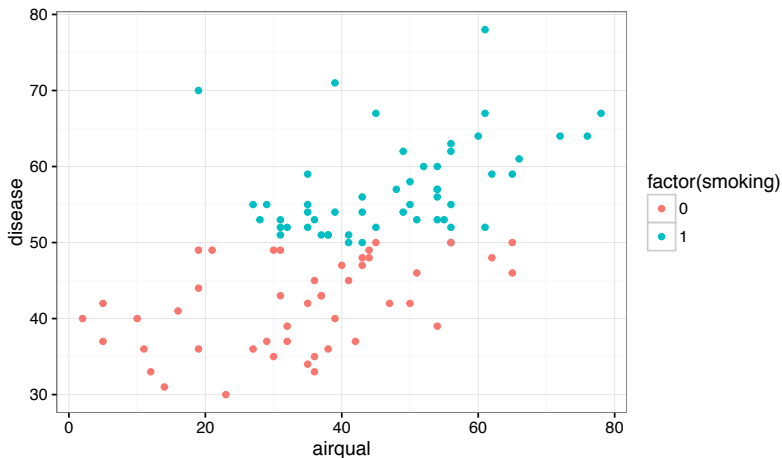
```
ggplot(dat, aes(crowding, disease)) + geom_point() +  
  geom_smooth(method="lm", se=FALSE)
```



What model syntax is implied by the above figure?

Lung Data Example: what is the model?

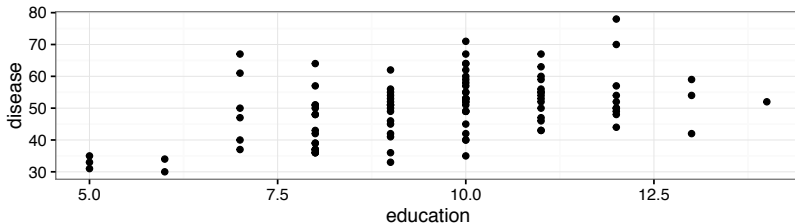
```
ggplot(dat, aes(airqual, disease, color=factor(smoking))) +  
  geom_point()
```



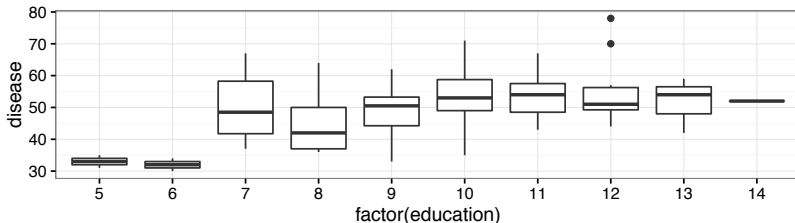
What is one possible model syntax implied by the above figure?

Which representation of education is better and why?

```
ggplot(dat, aes(education, disease)) + geom_point()
```



```
ggplot(dat, aes(factor(education), disease)) + geom_boxplot()
```



Lung Data Example

```
mlr1 <- lm(disease ~ crowding, data=dat)
kable(summary(mlr1)$coef, digits=2, format="latex")
```

| | Estimate | Std. Error | t value | Pr(> t) |
|-------------|----------|------------|---------|----------|
| (Intercept) | 12.99 | 3.48 | 3.74 | 0 |
| crowding | 1.51 | 0.14 | 10.83 | 0 |

```
mlr2 <- lm(disease ~ crowding + airqual, data=dat)
kable(summary(mlr2)$coef, digits=2, format="latex")
```

| | Estimate | Std. Error | t value | Pr(> t) |
|-------------|----------|------------|---------|----------|
| (Intercept) | 2.88 | 2.49 | 1.16 | 0.25 |
| crowding | 1.40 | 0.09 | 15.02 | 0.00 |
| airqual | 0.31 | 0.03 | 11.06 | 0.00 |

Why are the coefficients different?

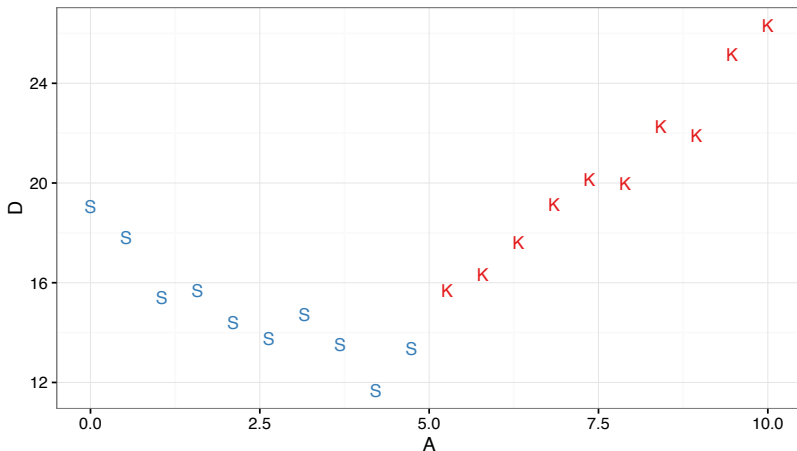
Lung Data Example

| | Estimate | Std. Error | t value | $\Pr(> t)$ |
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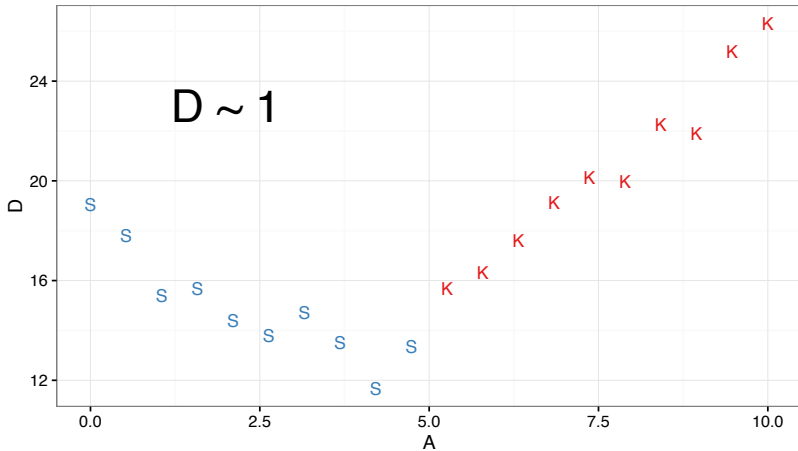
What are the interpretations of the coefficients?

Example data

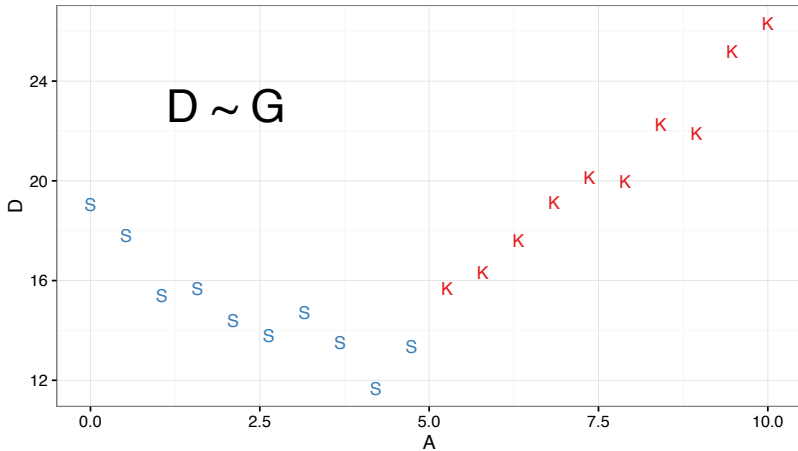
- D = a quantitative variable
- A = a quantitative variable
- G = a categorical variable with two levels, S and K



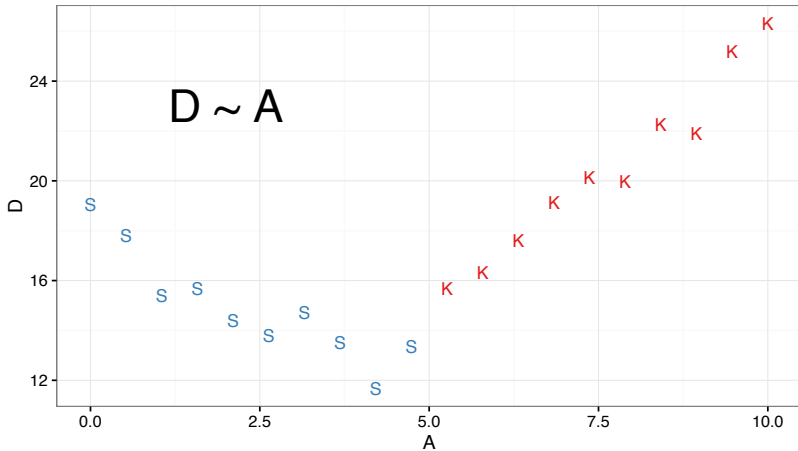
Draw the model...



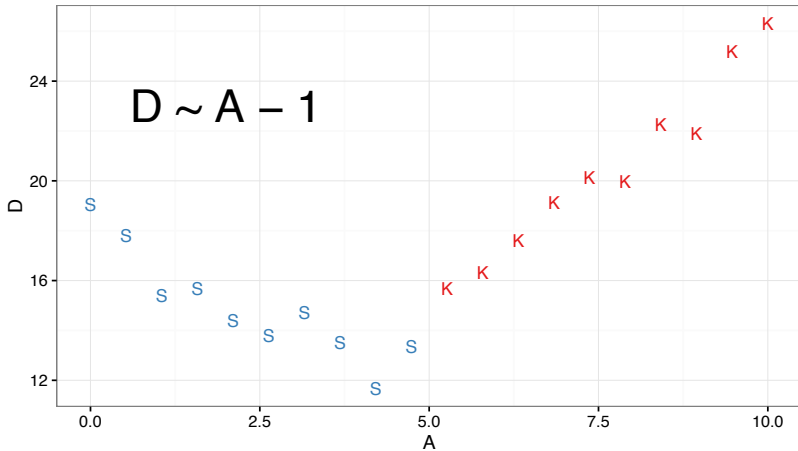
Draw the model...



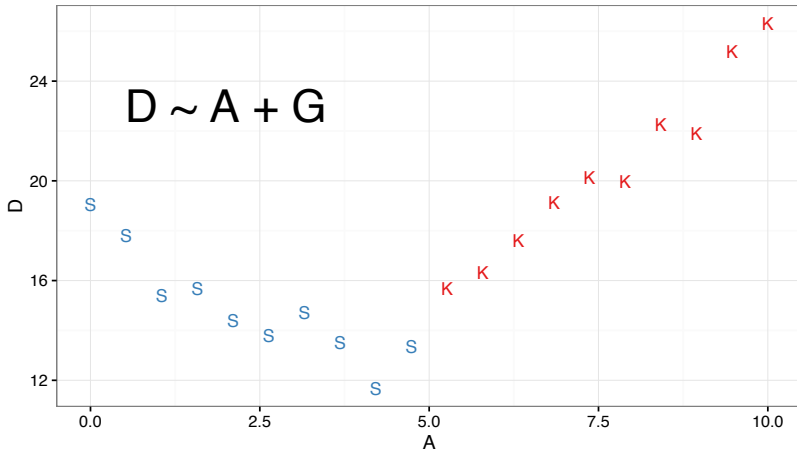
Draw the model...



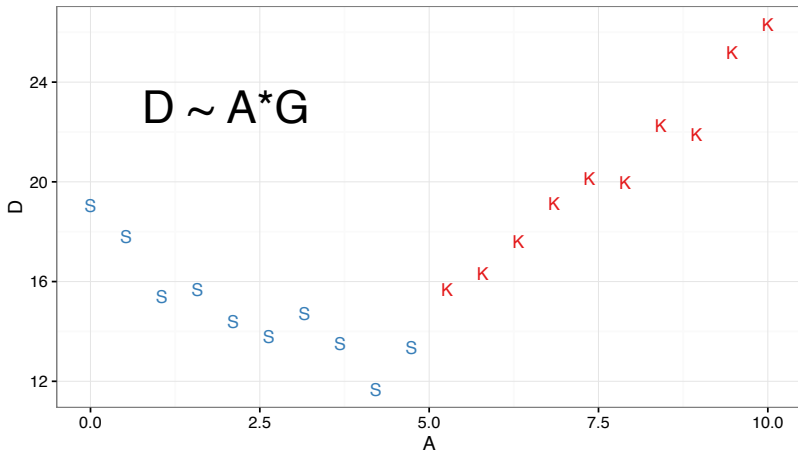
Draw the model...



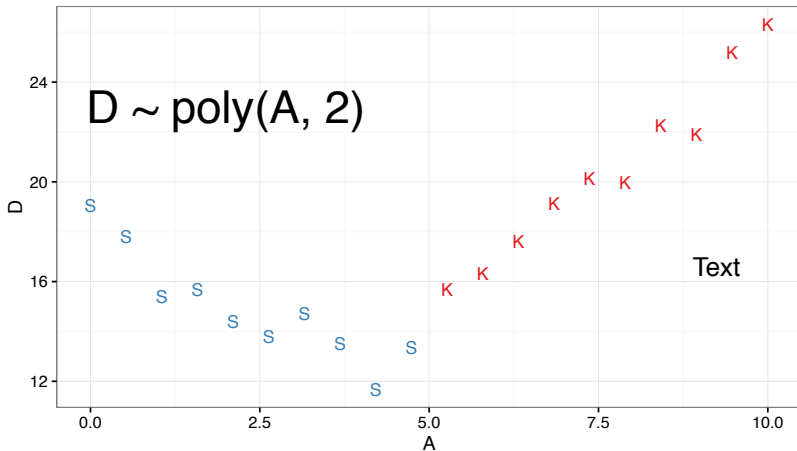
Draw the model...



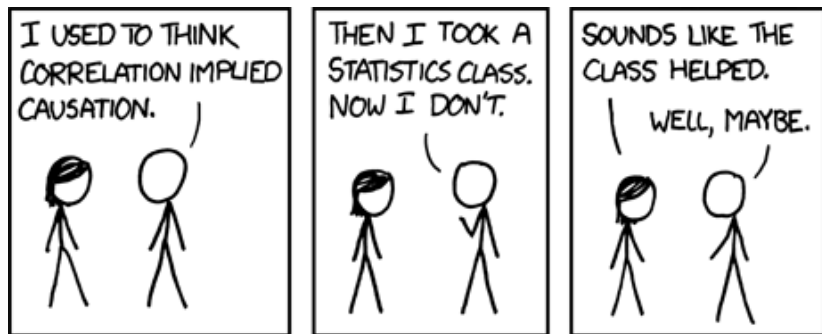
Draw the model...



Draw the model...



Parting wisdom



Up next: the mechanics and math of fitting models to data!

* Image credits: XKCD, <http://xkcd.com/552/>