

ealth

# Questions for this class

Introduction to regression

- Consider trying to answer the following kinds of questions:
  - To use the parents' heights to predict childrens' heights.
  - To try to find a parsimonious, easily described mean relationship between parent and children's heights.
  - To investigate the variation in childrens' heights that appears unrelated to parents' heights (residual variation).
  - To quantify what impact genotype information has beyond parental height in explaining child height.
  - To figure out how/whether and what assumptions are needed to generalize findings beyond the data in question.
  - Why do children of very tall parents tend to be tall, but a little shorter than their parents and why children of very short parents tend to be short, but a little taller than their parents? (This is a famous question called 'Regression to the mean'.)

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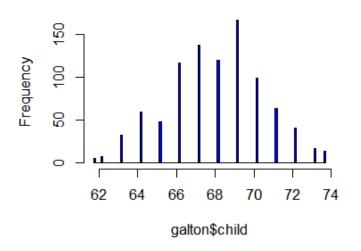
### Galton's Data

- Let's look at the data first, used by Francis Galton in 1885.
- · Galton was a statistician who invented the term and concepts of regression and correlation, founded the journal Biometrika, and was the cousin of Charles Darwin.
- You may need to run install.packages ("UsingR") if the UsingR library is not installed.
- Let's look at the marginal (parents disregarding children and children disregarding parents) distributions first.
  - Parent distribution is all heterosexual couples.
  - Correction for gender via multiplying female heights by 1.08.
  - Overplotting is an issue from discretization.

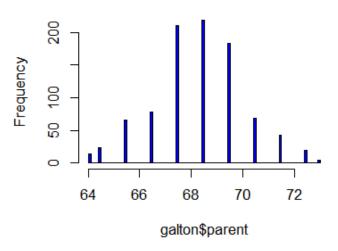
### Code

```
library(UsingR); data(galton)
par(mfrow=c(1,2))
hist(galton$child,col="blue",breaks=100)
hist(galton$parent,col="blue",breaks=100)
```

#### Histogram of galton\$child



#### Histogram of galton\$parent



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## Finding the middle via least squares

- · Consider only the children's heights.
  - How could one describe the "middle"?
  - One definition, let  $Y_i$  be the height of child i for  $i=1,\ldots,n=928$ , then define the middle as the value of  $\mu$  that minimizes

$$\sum_{i=1}^{n} (Y_i - \mu)^2$$

- · This is physical center of mass of the histrogram.
- You might have guessed that the answer  $\mu = \bar{X}$ .

### **Experiment**

Use R studio's manipulate to see what value of  $\mu$  minimizes the sum of the squared deviations.

```
library(manipulate)
myHist <- function(mu) {
  hist(galton$child,col="blue",breaks=100)
  lines(c(mu, mu), c(0, 150),col="red",lwd=5)
  mse <- mean((galton$child - mu)^2)
  text(63, 150, paste("mu = ", mu))
  text(63, 140, paste("MSE = ", round(mse, 2)))
}
manipulate(myHist(mu), mu = slider(62, 74, step = 0.5))</pre>
```

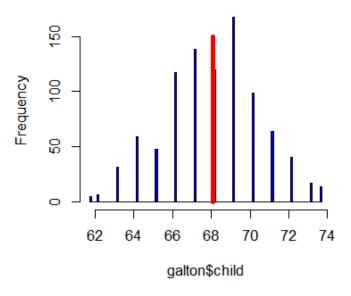
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## The least squares estimate is the empirical

### mean

```
hist(galton$child,col="blue",breaks=100)
meanChild <- mean(galton$child)
lines(rep(meanChild,100),seq(0,150,length=100),col="red",lwd=5)
```

#### Histogram of galton\$child



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### The math follows as:

$$\sum_{i=1}^{n} (Y_i - \mu)^2 = \sum_{i=1}^{n} (Y_i - \bar{Y} + \bar{Y} - \mu)^2$$

$$= \sum_{i=1}^{n} (Y_i - \bar{Y})^2 + 2 \sum_{i=1}^{n} (Y_i - \bar{Y})(\bar{Y} - \mu) + \sum_{i=1}^{n} (\bar{Y} - \mu)^2$$

$$= \sum_{i=1}^{n} (Y_i - \bar{Y})^2 + 2(\bar{Y} - \mu) \sum_{i=1}^{n} (Y_i - \bar{Y}) + \sum_{i=1}^{n} (\bar{Y} - \mu)^2$$

$$= \sum_{i=1}^{n} (Y_i - \bar{Y})^2 + 2(\bar{Y} - \mu)(\sum_{i=1}^{n} Y_i - n\bar{Y}) + \sum_{i=1}^{n} (\bar{Y} - \mu)^2$$

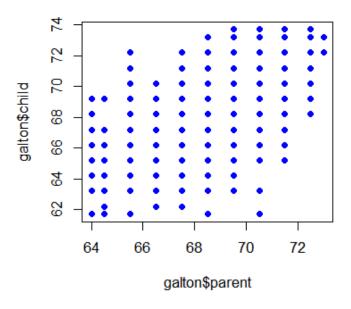
$$= \sum_{i=1}^{n} (Y_i - \bar{Y})^2 + \sum_{i=1}^{n} (\bar{Y} - \mu)^2$$

$$\geq \sum_{i=1}^{n} (Y_i - \bar{Y})^2$$

# Comparing childrens' heights and their

# parents' heights

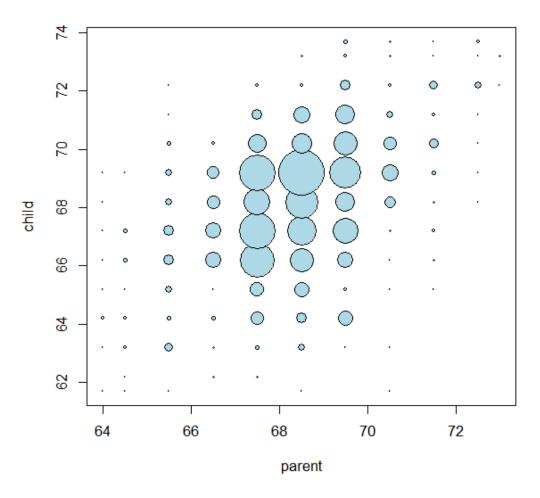
plot (galton\$parent, galton\$child, pch=19, col="blue")



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Size of point represents number of points at that (X, Y) combination (See the Rmd file for the code).

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# Regression through the origin

- Suppose that  $X_i$  are the parents' heights.
- · Consider picking the slope  $\beta$  that minimizes

$$\sum_{i=1}^{n} (Y_i - X_i \beta)^2$$

- This is exactly using the origin as a pivot point picking the line that minimizes the sum of the squared vertical distances of the points to the line
- · Use R studio's manipulate function to experiment
- · Subtract the means so that the origin is the mean of the parent and children's heights

myPlot <- function(beta) {</pre>

```
y <- galton$child - mean(galton$child)
  x <- galton$parent - mean(galton$parent)
  freqData <- as.data.frame(table(x, y))</pre>
  names(freqData) <- c("child", "parent", "freq")</pre>
  plot (
    as.numeric(as.vector(freqData$parent)),
    as.numeric(as.vector(freqData$child)),
    pch = 21, col = "black", bg = "lightblue",
    cex = .15 * freqData\$freq
    xlab = "parent",
    vlab = "child"
  abline (0, beta, lwd = 3)
  points (0, 0, cex = 2, pch = 19)
  mse \leftarrow mean((y - beta * x)^2)
  title(paste("beta = ", beta, "mse = ", round(mse, 3)))
manipulate (myPlot (beta), beta = slider (0.6, 1.2, step = 0.02))
```

### The solution

### In the next few lectures we'll talk about why this is the solution

```
lm(I(child - mean(child)) \sim I(parent - mean(parent)) - 1, data = galton)
```

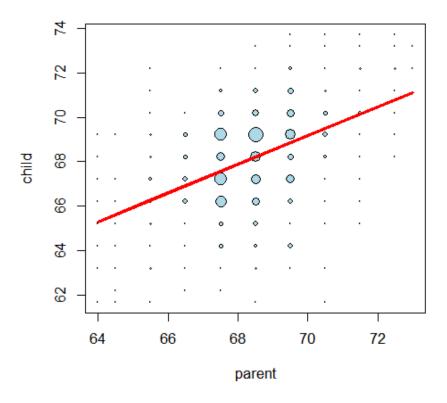
```
Call:
lm(formula = I(child - mean(child)) ~ I(parent - mean(parent)) -
    1, data = galton)

Coefficients:
I(parent - mean(parent))
    0.646
```

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# Visualizing the best fit line

### Size of points are frequencies at that X, Y combination



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