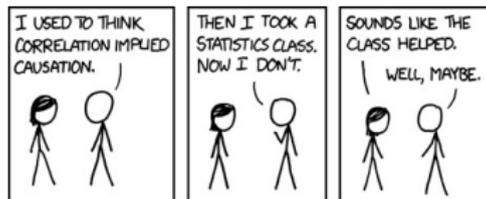


Correlation and Regression



<http://xkcd.com/552/>

How are X and Y Related

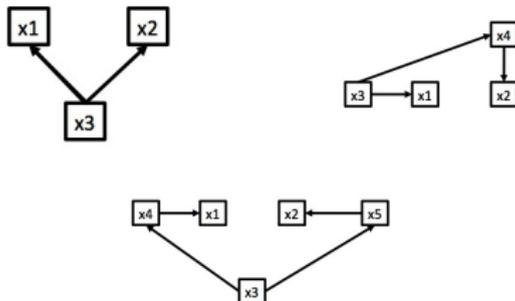


Causation (regression)
 $p(Y | X=x)$

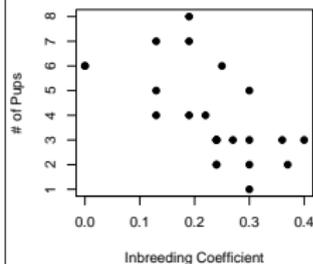


Correlation
 $p(Y=y, X=x)$

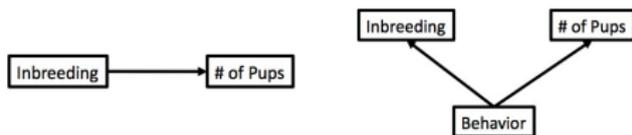
Correlation Can be Induced by Many Mechanisms



Example: Wolf Inbreeding and Litter Size



Example: Wolf Inbreeding and Litter Size



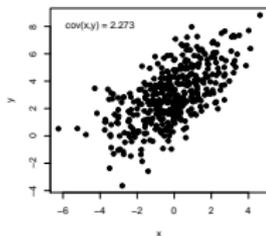
We don't know which is correct - or if another model is better. We can only examine *correlation*.

Covariance

Describes the relationship between two variables. Not scaled.

σ_{xy} = population level covariance, s_{xy} = covariance in your sample

$$\sigma_{XY} = \frac{\sum(X - \bar{X})(y - \bar{Y})}{n - 1}$$



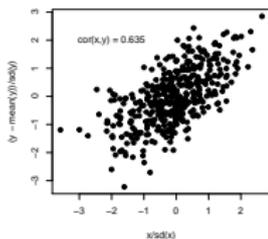
Pearson Correlation

Describes the relationship between two variables.

Scaled between -1 and 1.

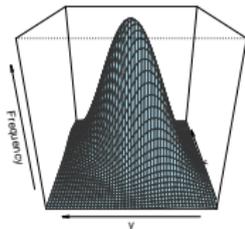
ρ_{xy} = population level correlation, r_{xy} = correlation in your sample

$$\rho_{xy} = \frac{\sigma_{xy}}{\sigma_x \sigma_y}$$



Assumptions of Pearson Correlation

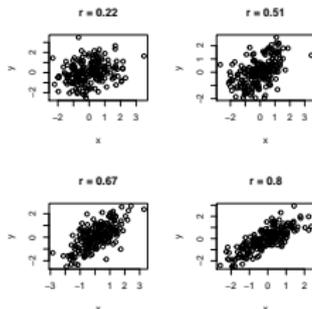
- ▶ Observations are from a **random sample**
- ▶ Each observation is **independent**
- ▶ X and Y are from a **Normal Distribution**



The meaning of r

Y is perfectly predicted by X if $r = -1$ or 1 .

r^2 = the porportion of variation in y explained by x



Testing if $r \neq 0$

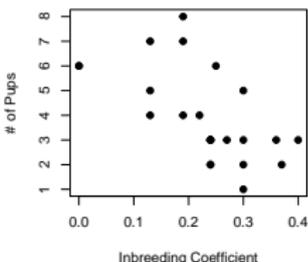
H_0 is $r=0$. H_a is $r \neq 0$.

Testing: $t = \frac{r}{SE_r}$ with $df=n-2$

WHY $n-2$?

$$SE_r = \sqrt{\frac{1-r^2}{n-2}}$$

Example: Wolf Inbreeding and Litter Size



Example: Wolf Inbreeding and Litter Size

```
cov(wolves)

##                               inbreeding.coefficient  pups
## inbreeding.coefficient           0.009922 -0.1136
## pups                             -0.113569  3.5199

cor(wolves)

##                               inbreeding.coefficient  pups
## inbreeding.coefficient           1.0000 -0.6077
## pups                             -0.6077  1.0000
```

Example: Wolf Inbreeding and Litter Size

```
with(wolves, cor.test(pups, inbreeding.coefficient))

##
## Pearson's product-moment correlation
##
## data: pups and inbreeding.coefficient
## t = -3.589, df = 22, p-value = 0.001633
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## -0.8120 -0.2707
## sample estimates:
## cor
## -0.6077
```

Exercise: Pufferfish Mimics & Predator Approaches

- ▶ Load up the pufferfish mimic data from W&S
- ▶ Plot the data
- ▶ Assess the correlation and covariance
- ▶ Assess H_0 .
- ▶ Challenge - Evaluate H_{a1} : the correlation is 1.



Exercise: Pufferfish Mimics & Predator Approaches

```
# get the correlation and se
puff_cor = cor(puffer)[1, 2]
se_puff_cor = sqrt((1 - puff_cor)/(nrow(puffer) - 2))

# t-test with difference from 1
t_puff <- (puff_cor - 1)/se_puff_cor
t_puff

## [1] -2.005

# 1 tailed, as > 1 is not possible
pt(t_puff, nrow(puffer) - 2)

## [1] 0.03013
```

Violating Assumptions?

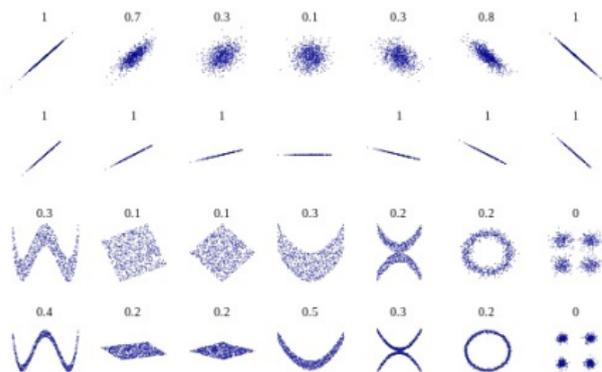
- ▶ Spearman's Correlation (rank based)
- ▶ Distance Based Correlation & Covariance (dcor)
- ▶ Maximum Information Coefficient (nonparametric)

All are lower in power for linear correlations

Spearman Correlation

1. Transform variables to ranks, i.e., 2, 3... (`rank()`)
2. Compute correlation using ranks as data
3. If $n \leq 100$, use Spearman Rank Correlation table
4. If $n > 100$, use t-test as in Pearson correlation

Distance Based Correlation, MIC, etc.

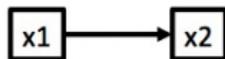


Least Squares Regression

$$y = ax + b$$

Then it's code in the data, give the keyboard a punch
Then cross-correlate and break for some lunch
Correlate, tabulate, process and screen
Program, printout, regress to the mean
-White Collar Holler by Nigel Russell

How are X and Y Related

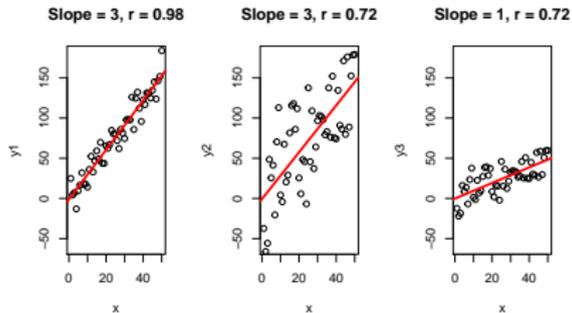


Causation (regression)
 $p(Y | X=x)$



Correlation
 $p(Y=y, X=x)$

Correlation v. Regression Coefficients

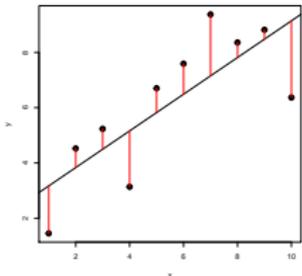


Basic Principles of Linear Regression

- ▶ Y is determined by X: $p(Y | X=x)$
- ▶ The relationship between X and Y is Linear
- ▶ The residuals of $Y = a + bx$ are normal distributed (i.e., $Y = a + bX + e$ where $e \sim N(0, \sigma)$)

Basic Principles of Least Squares Regression

$$\hat{Y} = a + bX - a = \text{intercept}, b = \text{slope}$$



$$\text{Minimize Residuals defined as } SS_{\text{residuals}} = \sum (Y_i - \hat{Y})^2$$

Solving for Slope

$$b = \frac{s_{xy}}{s_x^2} = \frac{\text{cov}(x,y)}{\text{var}(x)}$$
$$= r_{xy} \frac{s_y}{s_x}$$

Solving for Intercept

Least squares regression line always goes through the mean of X and Y

$$\bar{Y} = a + b\bar{X}$$

$$a = \bar{Y} - b\bar{X}$$

Fitting a Linear Model in R

```
wolf_lm <- lm(pups ~ inbreeding.coefficient, data=wolves)
```

```
wolf_lm
```

```
##  
## Call:  
## lm(formula = pups ~ inbreeding.coefficient, data = wolves)  
##  
## Coefficients:  
##      (Intercept)  inbreeding.coefficient  
##              6.57                -11.45
```

Extracting Coefficients from a LM

```
coef(wolf_lm)  
##      (Intercept) inbreeding.coefficient  
##              6.567                -11.447
```

```
coef(wolf_lm)[1]  
## (Intercept)  
##      6.567
```

Extracting Fitted Values from a LM

```
fitted(wolf_lm)
```

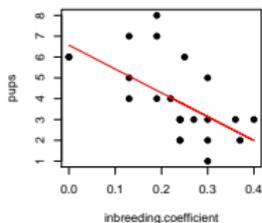
```
##      1      2      3      4      5      6      7      8      9     10  
## 6.567 6.567 5.079 5.079 5.079 4.392 4.392 4.392 3.706 3.820  
## 11     12     13     14     15     16     17     18     19     20  
## 3.820 3.820 3.820 3.820 3.820 3.477 3.133 3.133 3.133 3.133  
## 21     22     23     24  
## 2.446 1.989 2.332 4.049
```

```
coef(wolf_lm)[1] + coef(wolf_lm)[2]*0.25
```

```
## (Intercept)  
##      3.706
```

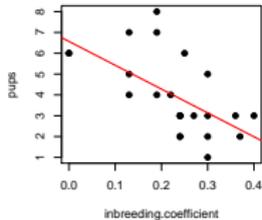
Plotting Fitted LMs

```
plot(pups ~ inbreeding.coefficient, data=wolves, pch=19)
matplot(wolves$inbreeding.coefficient, fitted(wolf_lm),
        add=T, lwd=2, col="red", type="l")
```



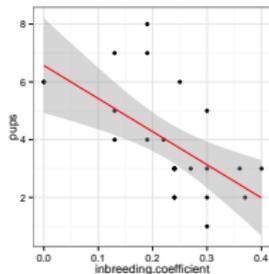
Plotting Fitted LMs

```
plot(pups ~ inbreeding.coefficient, data=wolves, pch=19)
abline(wolf_lm, col="red", lwd=2)
```



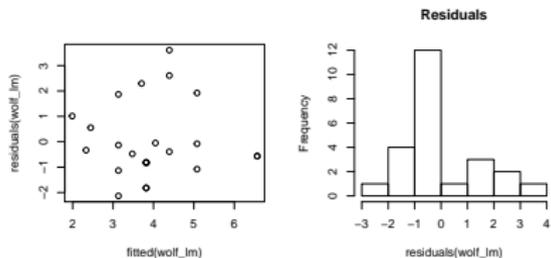
Ggplot2 and LMs

```
ggplot(data=wolves, aes(y=pups, x=inbreeding.coefficient)) +
  geom_point() +
  theme_bw() +
  stat_smooth(method="lm", color="red")
```



Checking Residuals

```
par(mfrow=c(1,2))
plot(fitted(wolf_lm), residuals(wolf_lm))
#
hist(residuals(wolf_lm), main="Residuals")
```



Exercise: Pufferfish Mimics & Predator Approaches

- ▶ Fit the pufferfish data
- ▶ Visualize the linear fit
- ▶ Examine whether there is any relationship between fitted values, residual values, and treatment

