Parameter Contrasts: Regression Framework JH notes on regression

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Introduction to parameter-contrasts

- We started the course by talking about the case where there were no determinants, i.e., no subpopulations \rightarrow there was one global parameter (μ , π , λ).
- Now we concern ourselves with determinants of the global parameter. For example:
 - \blacktriangleright μ_{north} VS. μ_{south}
 - \blacktriangleright π_{north} VS. π_{south}
 - $ightharpoonup \lambda_{north}$ vs. λ_{south}
- Today we introduce population parameter <u>contrasts</u> in a regression framework

Why regression for parameter-contrasts?

- Why do we start in a regression framework (as opposed to two-sample inference in B&M and AAO)?
- Parameter contrasts are a special case of regression
- Approach taken in Miettinen, Clayton in Hills, Rothman and Greenland, baby Rothman

What is regression?

- How parameters relate to its determinants
- How to link the parameters between the different populations through generic equations, that looks like a regression equation.
- Then once you get data, you can actually fit or get your best estimates of those parameters

Linear regression: The Concept

A regression model is said to be linear when it is of the form

$$\mu = \mu_0 + \sum_{j=1}^{p} \beta_j X_j$$

= $\mu_0 + \beta_1 X_1 + \beta_1 X_1 + \dots + \beta_p X_p$

• Which means that the value of the mean (μ) is viewed as a linear combination of the parameters $\mu_0, \beta_1, \beta_2, \ldots, \beta_p$, the coefficients of the linear combination being the realizations for the X's

Linear regression: Example

- Consider the depths of the ocean example
- lacktriangle Here, μ designates the true mean depth of the ocean
- For this parameter, one might consider the determinant
 - ► X₁: South hemisphere, represented by an indicator variable

$$X_1 = \begin{cases} 1 & \text{if Southern hemisphere} \\ 0 & \text{if Northern hemisphere} \end{cases}$$

Linear regression: Example

■ The model might be taken as

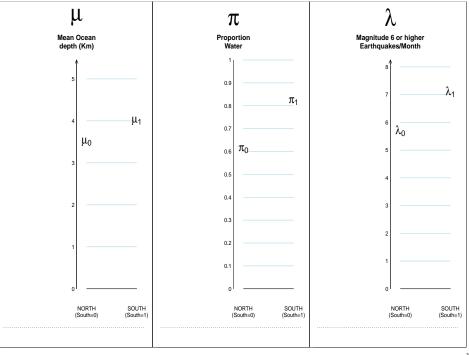
$$\mu = \mu_0 + \beta_1 X_1$$

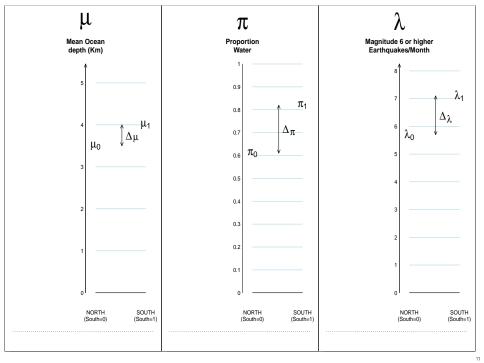
and provides the mean depth of the ocean $\underline{\text{given}}\ X_1$

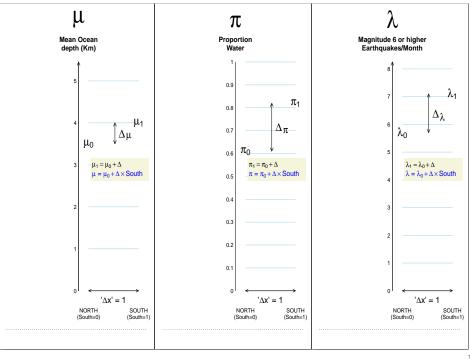
- The mean depth of the ocean μ is a linear combination of μ_0 and β_1
- If we had an infinite amount of data, the mean depth of the ocean would be determined by hemisphere:

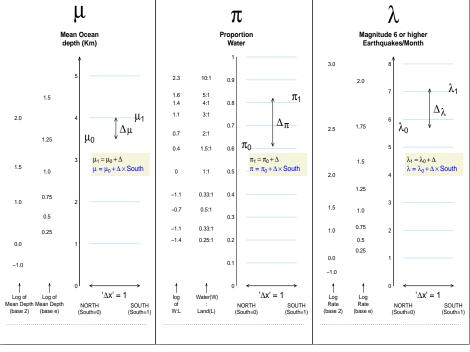
$$\mu = \begin{cases} \mu_0 + \beta_1 & \text{if Southern hemisphere} \\ \mu_0 & \text{if Northern Hemisphere} \end{cases}$$

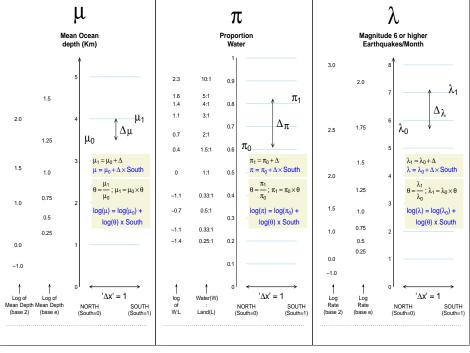
Regression equations when the truth is known

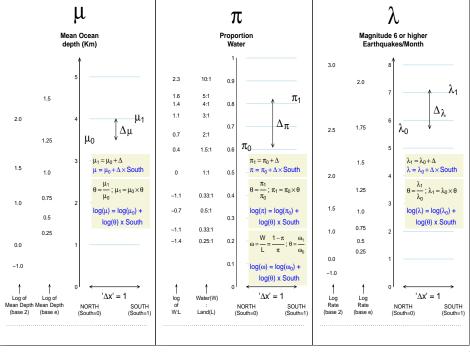


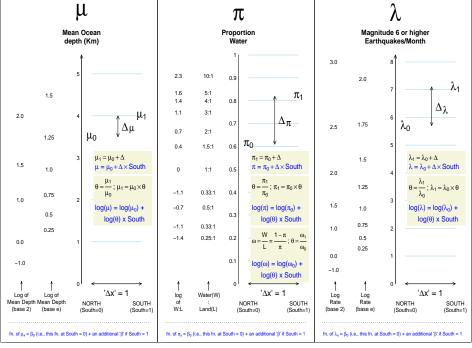










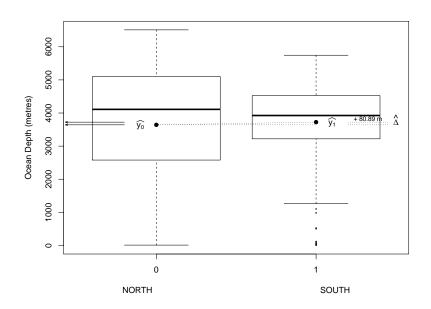


Fitting the regression equation with our sample data

Depths of the ocean: North vs. South Hemisphere

```
# load function to get depths
source("https://github.com/sahirbhatnagar/EPIB607/raw/master/
exercises/water/automate water task.R")
# get 1000 depths
set.seed(222333444)
depths <- automate_water_task(index = sample(1:50000, 1000),
        student id = 222333444, type = "depth")
# separate by north and south hemisphere
depths north <- depths[which(depths$lat>0),]
depths south <- depths[which(depths$lat<0),]
# restrict sample to 200 (at random)
depths north <- depths north[sample(1:nrow(depths north), 200), ]
depths south <- depths south[sample(1:nrow(depths south), 200), ]
# add indicator variable
depths north$South <- 0
depths south$South <- 1
# combine data
depths <- rbind(depths north, depths south)
head(depths)
# calculate mean and sd by hemisphere
means <- aggregate(x = depths, by = list(depths$South), FUN = "mean")$alt
sds <- aggregate(x = depths, by = list(depths$South), FUN = "sd")$alt
```

Depths of the ocean: North vs. South Hemisphere



Standard error of the mean difference

To perform inference we first need to calculate the SE of the mean difference given by:

$$SE_{\bar{y_1} - \bar{y_0}} = \sqrt{\frac{s_0^2}{n_0} + \frac{s_1^2}{n_1}} \tag{1}$$

```
n0 <- nrow(depths_north)
n1 <- nrow(depths_south)

mean0 <- mean(depths_north$alt)
mean1 <- mean(depths_south$alt)

var0 <- var(depths_north$alt)
var1 <- var(depths_south$alt)

(SEM <- sqrt(var0/n0 + var1/n1))
## [1] 157.565</pre>
```

95% Confidence Interval for the Mean Difference

We can then calculate a 95% CI for the mean difference given by:

$$(\bar{y_1} - \bar{y_0}) \pm t^*_{(n_0 + n_1 - 2)} \times SE_{\bar{y_1} - \bar{y_0}}$$
 (2)

Parameter contrasts with regression

Using the lm function in R:

```
# regression. lm assumes equal variances
fit <- lm(alt ~ South, data = depths)</pre>
summary(fit)
##
## Call:
## lm(formula = alt ~ South, data = depths)
##
## Residuals:
      Min 10 Median 30 Max
##
## -3722.0 -608.5 401.5 1200.4 2867.9
##
## Coefficients:
             Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 3643.08 111.42 32.698 <2e-16 ***
## South 80.88 157.56 0.513 0.608
##
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1576 on 398 degrees of freedom
## Multiple R-squared: 0.0006617, ^ IAdjusted R-squared: -0.001849
## F-statistic: 0.2635 on 1 and 398 DF, p-value: 0.608
```

Confidence interval from regression fit

confint(fit)

```
## 2.5 % 97.5 %
## (Intercept) 3424.0440 3862.1160
## South -228.8787 390.6487
```

Unequal variances using stats::t.test

stats::t.test assumes unequal variances by default:

```
t.test(alt ~ South, data = depths, var.equal = FALSE)
##
## ^^IWelch Two Sample t-test
##
## data: alt by South
## t = -0.51334, df = 349.62, p-value = 0.608
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -390.7795 229.0095
## sample estimates:
## mean in group 0 mean in group 1
         3643.080 3723.965
##
(mean0 - mean1) + qt(c(0.025, 0.975), df = 349.61783) * SEM
## [1] -390.7795 229.0095
```

Equal variances using stats::t.test

We can specify equal variance assumption in stats::t.test:

```
t.test(alt ~ South, data = depths, var.equal = TRUE)
##
## ^^ITwo Sample t-test
##
## data: alt by South
## t = -0.51334, df = 398, p-value = 0.608
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -390.6487 228.8787
## sample estimates:
## mean in group 0 mean in group 1
##
         3643.080 3723.965
(mean0 - mean1) + qt(c(0.025, 0.975), df = n0 + n1 - 2) * SEM
## [1] -390.6487 228.8787
```