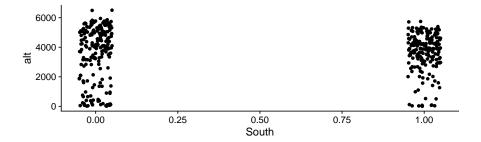
1 Mean depth of the ocean

```
head(depths)
            X
                     lon
                               lat alt water South
## 41995 41995 -87.21236 59.290367 190
## 11151 11151 -122.33034 5.554558 4167
## 43640 43640 -148.54790 36.237464 5447
## 8615
         8615 -24.92364 21.625967 5063
                                                  0
## 8126
         8126 177.18458 13.880370 5634
                                                  0
## 16548 16548
                48.88215 3.229250 3691
                                                  0
dim(depths)
## [1] 400 6
fit <- lm(alt ~ 1, data = depths)</pre>
summary(fit)
##
## Call:
## lm(formula = alt ~ 1, data = depths)
## Residuals:
      Min
               1Q Median
                               3Q
                                      Max
## -3681.5 -584.8 405.5 1197.2 2827.5
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 3683.52
                            78.71
                                     46.8 <2e-16 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 1574 on 399 degrees of freedom
```

2 Mean depth of the ocean in northern and southern hemisphere



```
fit <- lm(alt ~ South, data = depths)</pre>
summary(fit)
##
## Call:
## lm(formula = alt ~ South, data = depths)
## Residuals:
      Min
               1Q Median
                                      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 ***
                 80.88
                           157.56 0.513
## ---
## 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, Adjusted R-squared: -0.001849
## F-statistic: 0.2635 on 1 and 398 DF, p-value: 0.608
t.test(alt ~ South, data = depths, var.equal = TRUE)
##
   Two 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
```

3 Ratio depth of the ocean in northern and southern hemisphere

```
# note: we are now using glm
fit <- glm(alt ~ South, data = depths, family = gaussian(link=log))</pre>
summary(fit)
##
## Call:
## glm(formula = alt ~ South, family = gaussian(link = log), data = depths)
## Deviance Residuals:
      Min
                1Q Median
                                  3Q
                                          Max
## -3722.0
            -608.5
                    401.5
                             1200.4
                                       2867.9
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 8.20058
                          0.03058 268.144
                                            <2e-16 ***
               0.02196
## South
                          0.04278 0.513
                                             0.608
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for gaussian family taken to be 2482673)
##
      Null deviance: 988758010 on 399 degrees of freedom
## Residual deviance: 988103771 on 398 degrees of freedom
## AIC: 7029.1
## Number of Fisher Scoring iterations: 5
```

4 Student drinking

```
fit <- lm(drinks ~ gender, data = drinks)
summary(fit)
##
## Call:
## lm(formula = drinks ~ gender, data = drinks)
## Residuals:
      Min
              1Q Median
                             30
## -5.5185 -1.7947 -0.2947 1.4815 9.4815
## Coefficients:
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 4.2947
                        0.2837 15.138 < 2e-16 ***
## gender
               2.2238
                       0.4182 5.318 3.2e-07 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 2.765 on 174 degrees of freedom
## Multiple R-squared: 0.1398, Adjusted R-squared: 0.1348
## F-statistic: 28.28 on 1 and 174 DF, p-value: 3.197e-07
fit <- glm(drinks ~ gender, data = drinks, family = gaussian(link=log))</pre>
summary(fit)
##
## glm(formula = drinks ~ gender, family = gaussian(link = log),
      data = drinks)
## Deviance Residuals:
      Min
           1Q Median
## -5.5185 -1.7947 -0.2947 1.4815 9.4815
## Coefficients:
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 1.45739 0.06606 22.062 < 2e-16 ***
              ## gender
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for gaussian family taken to be 7.646385)
##
      Null deviance: 1546.7 on 175 degrees of freedom
## Residual deviance: 1330.5 on 174 degrees of freedom
## AIC: 861.48
## Number of Fisher Scoring iterations: 5
```